<400% 0317 ggdaagdgtt gtgdagagd	19
H210 + 2318 H211 + 19	
HCC12 + DNA HCC13 + Homo sapiens	
<pre><d::00 -="" <d::223="" binding="" cyclin="" e="" pre="" ribbzyme="" site<=""></d::00></pre>	
k400 - 1318 agagtesata gecagetgg	19
<210x 2319	
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<220 - <223 Cyclin F ribozyme binding site	
-:400>-2319	
oggogtggto cactgtagg	19
<pre>&lt;210 % 2320 <pre>&lt;211 % 19</pre></pre>	
<pre><c12 +="" dna<="" pre=""></c12></pre>	
-213 Homo sapiens	
<pre>&lt;0008 &lt;0008</pre>	
-:4008-2320 gtodactgta-ggtgtgcca	19
<2108 2321	
F0118-19 F0108-DNA	
<pre>% Company </pre>	
+(20.0)8	
+2130 Cyclin F ribozyme binding site	
+14001+ 1711	<b>,</b>
qccaag*qtt totgttato	19
+0.10+ 1.322 +0.11+ 1.9	
H12120 DNA	

<pre>&lt;:213   Homo sapiens</pre>	
RUBBA - Gyplin F ribozyme binding site	
H400 + 0322 possytyttt otgitatoo	19
<pre>%210% 2303 %211 * 19 %212 * DNA %213 * Home sapiens</pre>	
+07.0> <0003	
<400 > 2323 caaqtgttto tgttatoot	19
<pre>K310 % 3334 K311 * 19 K312 * DNA K313 * Homo sapiens</pre>	
<2220° <223 - Cyclin F ribozyme binding site	
<pre>&lt;:400 - 2324 tgtttctgtt atcctacaa</pre>	19
+0210 + 0325 +0211 + 19 +0012 + DNA +0013 + Homo sapiens	
+12005 +1223 + Cyclin F ribozyme binding site	
<pre></pre> <pre><pre><pre></pre> <pre><pre>qtttetatta tootacaaa</pre></pre></pre></pre>	19
+00108 0326 +00118 19 +00118 DNA +00138 Homo sapiens	
+:210:- -:2:30- Cyclin F ribozyme binding site	
<pre>&lt;:400 - 2326 ttctgttatc_ctacaaagc</pre>	19

<pre>K210 \ 2327 K211 + 1# K212 + DNA K213 + Homo sapiens</pre>	
<pre><pre><pre><pre>&lt;223 - Cyclin F ribozyme binding site</pre></pre></pre></pre>	
H400 - 2327 tyttatoota caaagogaa	19
<pre>#210 + 0328 #211 + 19 #212</pre>	
<pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre>&lt;</pre></pre>	
-:4005-2328 qoqaagaata aggaggagg	19
<pre>%010% 2329 %011% 19 %012% DNA %013% Homo sapiens</pre>	
+CCO+ +CCC3+ Cyclin F ribozyme binding site	
<400% 0309 cotqaccato ttgagtoto	19
\$210 \ 2330 \2212 19 \2212 DMA \$210 Hemo sapiens	
-MANGE Cyclin F ribozyme binding site	
+40f5 [336] tgadbatoft gagtotobo	19
+0108 [331 +0118 [14 +0118 DNA +0213 Homo sapiens	
+:220×	

Hamas Cyclin F ribozyme binding site	
<pre>At Minga its tecoogaag</pre>	19
+C10 + 2332 +C11 + 19 +C12 + DNA +C13 + Homo sapiens	
<pre><pre></pre></pre>	
ALL3 - Cyclin F ribozyme binding site	
-:400 + 2332 	19
<pre>%210 * 2333 %211 * 19 %212 * DNA %213 * Homo sapiens</pre>	
<pre>*D20* KD23 + Gydlin F ribezyme binding site</pre>	
k400 - 2333 adatatgeto tittoacato	19
+210 + 2334 +211 + 19 +212 + DNA +213 + Homo sapiens	
k2208 K223 Cyclin F ribozyme binding site	
<pre>c400% 2034 atquactett teacatest</pre>	19
+D10+ 1335 +D11+ 19 +D13+ DNA +D13+ Homo sapiens	
-Mile- -Mile-Cyclin F ribozyme binding site	
<pre>04000 2335 igtwoto:tt dacatootg</pre>	19
-0210 + 2036 -0211 > 19	

<pre>0313 &gt; DNA 0313 &gt; Homo sapiens</pre>	
<pre>000000 000000000000000000000000000000</pre>	
H400+ 1336 Atgetettte acatectga	19
+1210 + 2337 +1211 + 13 +1212 + DMA +1213 + Homo sapiens	
<pre>&lt;220 - &lt;223 - Cyclin F ribozyme binding site</pre>	
<400> 2337 otttoacato otgaaatgg	19
<210 > 2338 <211 > 19 <212 > DNA <213 > Homo sapiens	
<220> <223× Oyclin F ribozyme binding site	
<400% 2338 qaaatgqott totgtagag	19
<pre>\$210 \( 0339 \) &lt;211 \( 19 \) &lt;212 \( \text{PMA} \) &lt;13 \( \text{Homo sapiens} \)</pre>	
-0.00 s -0.03 - Cyclin F ribozyme binding site	
H400% 1934 Waatggett: otgtagagg	19
+110:- 234:- +111:- 19 +111:- DNA +113:- Homo sapiens	
<pre>+210: &lt;213: Cyclin F ribozyme binding site</pre>	
+.400:> ∠340	

aatggettte tgtagagga	19
+3210 + 3341 +3211 + 13 +3212 + LNA +3213 + Homo sapiens	
+:220 + -: -:223 - Cyclin F ribozyme binding site	
-:400 - 2341 -potetotata gaggacato	19
+0.100 2342 +0.11> 19 +0.12> DNA +0.13> Homo sapiens	
<pre>&lt;100&gt; &lt;1003 Cyclin F ribozyme binding site</pre>	
<400 · 0340 agaggacato otggoogto	19
K210 + 2343 K211 + 19 K212 + DMA K213 + Homo sapiens	
+:2205 +:223: Cyclin F ribozyme binding site	
k(4005-2343 cotggoogto ogagotgta	19
<pre>%10.08 0.344 %11.18 1.9 %21.08 EMA %21.38 Homo sapiens</pre>	
+02008 +023> Cyclin F ribozyme binding site	
<pre>-:400:- 0:444 ccgagetita cacteccag</pre>	19
+0010+ 1345 +0010+ 13 +0010+ FMA +0010+ FMA +0013> Home sapiens	

<pre>+:220 + +:223 + Cyclin F ribozyme binding site</pre>	
+:400 + 2345	1.0
otqtacaoto ccagotgaa	19
+0010 × 1346	
H011 × 19	
HARLEY DNA	
HD13 - Homo sapiens	
+:220 +	
+:223% Cyclin F ribozyme binding site	
+:400 + 2346	
gtgobagott ccaggagot	19
, , , , , a garage a	
<210× 2347	
<2118-19	
<pre><!--DIL--> DNA</pre>	
+:213× Homo sapiens	
<1110 >	
<2223 Cyclin F ribozyme binding site	
<400> 0347	1.0
tqqqagqtta caggagqtg	19
<210 × 2348	
kD11> 19	
HD12> DNA	
<pre>&amp;213&gt; Homo sapiens</pre>	
+(2208)	
<pre><d:d3> Cyclin F ribozyme binding site</d:d3></pre>	
+(400% 1349	19
totggoogto tocagggaa	13
+:2100+ 2349	
(C11) 19	
+212+ DNA	
+2130 Homo sapiens	
*(10.0)*	
+3733+ Cyclin F ribozyme binding site	
+(400)+ 2344	10
tagenagticta cagggaacc	19
0.210 > 2350	

<pre>#211** 19 #2112 * DUA #2113 * H mo sapiens</pre>	
+0.20 + + Oyolin F ribozyme binding site	
H400 + 1750 Solyaaqoto tttgaaagg	19
+0210 + 2351 +0211 + 19 +0212 + DNA +0213 * Homo sapiens	
<220> <223 Cyclin F ribozyme binding site	
<pre>&lt;400&gt; 2351 tgaagetett tgaaaggge</pre>	19
<pre>SC10 &gt; C352 SC211 &gt; 19 CC12 &gt; DNA CC213 &gt; Homo sapiens</pre>	
<pre><i220 +="" -="" <i223="" binding="" cyclin="" f="" pre="" riboxyme="" site<=""></i220></pre>	
-:4υθ - 2352 gaaqetettt gaaaggget	19
<pre>#0010% 2353 #0010% 19 #0010% DNA #0013% Homo sapiens</pre>	
RC210 RC230- Cyclin F ribozyme binding site	
+:400:+ 1353 aadqugaatt togaagotg	19
+00160+ 1054 +0110+ 19 +0110+ DNA +0110+ Homo sapiens	
-0.00- -0223> Cyclin F ribozyme binding site	

ा400 + 2354 अवद्याप्रवास tt ogaagotgo	19
00010 + 0355 0011 + 19 0010 + DNA 0013 + Humo sapiens	
<pre><pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><!--</td--><td></td></pre></pre></pre></pre>	
्र400 - 2356 अववृत्रaatite gaagetget	19
<pre>4010 + 2356 4211 &gt; 19 4010 &gt; DNA 4013 &gt; Homo sapiens</pre>	
<pre>&lt;2000&gt; &lt;2003   Oyolin F ribozyme binding site</pre>	
-:400+ ::356 gotaggoata gootacoto	19
+0.10 x 2357 +0.11 x 19 +0.210 + DNA +0.213 + Homo sapiens	
-CDDG - -CDDB- Cyclin F ribozyme binding site	
-04000-1357 quatagusta cetetacaa	19
+010+ 235+ +0.11* 14 +0.10* PNA +0.13* Homo sapiens	
<pre>:C100: :C130: Cyclin F ribozyme binding site</pre>	
-04000- 2358 agostassts tacaatgaa	19
+00100 L+53 +00100 L+53 +00100 DMA +00130 Home samiens	

Hanner of the Ha	
-(40) 2359 Dotabotota caatgaagg	19
<pre>%210 * 2360 %211 * 19 %212 * DNA %213 * Homo sapiens</pre>	
edlo. edlo. Syslin F ribozyme binding site	
<pre>&lt;400 + 2360 aaqqcotqto tgtgtotga</pre>	1,9
<pre>#010 + 0361 +011 &gt; 19 #010 + DMA #013 &gt; Homo sapiens</pre>	
<pre><pre><pre><pre>&lt;220 *</pre> <pre>&lt;223 &gt; Cyclin F ribozyme binding site</pre></pre></pre></pre>	
<pre>c400x 2361 tgtotgtgto tgatgaggo</pre>	19
<pre>*C10 &gt; C36C *C11 &gt; 19 *C12 &gt; DNA *C13 &gt; Homo sapiens</pre>	
<pre>&lt;2008 &lt;2008 cyclin F ribozyme binding site</pre>	
-0400 = 0360 tgaaggooto togottott	19
+1110 + 1.363 +2111 + 19 +1111 + DNA +1112 + Homo sapiens	
CLICO: CLICK: Cyclin F ribozyme binding site	
::400:- 2363 aaggeetete gettettea	19

+:210 + 2364 :211 + 14 -:211 + DMA -:213 + Himb sapiens	
<pre>0326* 0323* Cyclin F ribozyme binding site</pre>	
H400 - 1364 destatoratt officaglet	19
+1210 × 1365 +1211 + 19 +1212 + DMA +1213 + Homo sapiens	
<pre>&lt;220* &lt;223 - Cyclin F ribezyme binding site</pre>	
<pre>&lt;400 &gt; 2365 etchettett cagtotogo</pre>	19
<pre>%210% 2366 %211% 19 %212% DNA %213% Homo sapiens</pre>	
<pre>s:210 + +223 - Cyclin F ribczyme binding site</pre>	
+4000-0366 togottotto agtotogot	19
\$2105 2367 \$2115 19 \$2128 DMA \$2135 Home sapiens	
<pre>87.08 87.38 Cyclin F ribozyme binding site</pre>	
<pre>+400% 1367 thefteagte tegetgage</pre>	19
+02160 2068 +0118 19 +0110 DNA +0110 Homo sapiens	
+C MC+ +CZ23> Cyclin F ribozyme binding site	

-(4))+ 2363 ottoagnoto gotgagogg	19
H210 + H369 H211 + 19 H212 + DNA H213 + Homo sapiens	
<pre>+:220 + +:223 + Cyclin F ribozyme binding site</pre>	
ा400× 1369 ब्रह्माल्यक्वाक्टर्स teatetgge	19
<pre>&lt;010 - 2370 &lt;011 - 19 &lt;010 DNA</pre>	
<pre><pre><pre><pre></pre> <pre><pre></pre> <pre></pre> <pre><pre></pre> <pre></pre> <pre></pre> <pre><pre></pre> <pre></pre> <pre></pre></pre></pre></pre></pre></pre></pre>	
+:400 + 2370 Degeacetts catetgget	19
+0210 + 2371 +0211 + 19 +0212 * DNA +0213 + Homo sapiens	
<pre>&lt;:2205 ::2235 Cyclin F ribozyme binding site</pre>	
+:400 = 2371 pacapotito atotgata	19
+2108 2372 +2118 10 +2128 DNA +2138 Homo sapiens	
<pre>%2208 %223% Cyclin F ribczyme binding site</pre>	
<pre>+:400:- 1372 additionate tggetette</pre>	19
+010+ 1373 +211+ 19 +212+ DNA	

Homo sapiens	
+0.120 +	
+MBB+ Cyclin F ribozyme binding site	
+1400 + 2373	
pathologicate theateoge	19
+1210 + 2374 +1211 + 19	
HOIL DNA	
+Cll3+ Homo sapiens	
+12100 +	
+223 - Cyclin F ribozyme binding site	
+:400 × 0.374	
totggetett categgese	19
<2108 2375 R2118 19	
*:212 DNA	
<pre>*:213 * Homo sapiens</pre>	
x(220)x	
<pre>&lt;223&gt; Cyclin F ribozyme binding site</pre>	
<:4008_2375	
etggetette ateegeest	19
H2108 2376 H2118 19	
KO10× DNA	
<pre><pre><pre><pre>#2135 Homo sapiens</pre></pre></pre></pre>	
+ 200 s	
+213 Oyolin F ribozyme binding site	
<4008 . 376	1.0
getatteate egecotoeg	19
+D10+ D377	
H2115 19 H2115 DNA	
+21 % Homo sapiens	
+/2L/**-	
+22% Cyclin F ribozyme binding site	
+(400)> 2377	
atdegecete egtggtegg	19

+1210	
<pre><i220 -="" <i223="" binding="" cyclin="" f="" pre="" ribozyme="" site<=""></i220></pre>	
H400 × 2378 Stocatgato ggtgagogg	19
<pre>HD10 &gt; 2379 HD11 + 19 HD12 &gt; DNA <d13> Homo sapiens</d13></pre>	
<2208 <2223 Cyclin F ribozyme binding site	
ार्थ00 र 2379 व्यवक्ष्युर्वेष्ट्रपूर्व dacgagage	19
+010 × 0380 +011 × 19 +0010 × DNA +0013 × Homo sapiens	
+22008 +22008 Cyclin F ribozyme binding site	
k:400:- 0380 geogtgatta acgagaged	19
<pre>&lt;210% 2381 &lt;2210% 19 &lt;2110% DNA &lt;2110% Homo sapiens</pre>	
-CTMS Cyclin F ribozyme binding site	
H400H 0381 ogađagosto agggoagag	19
+0100+0382 +0211+19 +0017+ DNA +0013+ Homo sapiens	
+:22:0:>	

<pre>&lt;223 - Cyclin F ribozyme binding site</pre>	
H400+ U382 bayayganto acaaagcat	19
+1210 + 2383 +1211 > 19 +1212 + DNA	
AMMIN Homo sapiens	
+020 + 4223 * Cyclin F ribozyme binding site	
<pre>&lt;400 + 2383 adaaaagdato datattgda</pre>	19
H210 + 2384 H211 + 19	
<pre>%212% DNA %213   Homo sapiens</pre>	
+0000+ +0003+ Cyclin F ribozyme binding site	
+400+ 2384 agratocata tigoacigo	19
HILLON 1385	
HD115 19 HD105 DNA	
+2213> Humo sapiens	
<pre>&lt;:210&gt; &lt;223&gt; Cyclin F ribozyme binding site</pre>	
+:400's 2385	
catocatatt gcactgctt	19
<0108 . 386 <01108 19	
STOLING DEA	
+0013 · Homo sapiens	
Hadisələr	
+1130 Cyclin F ribozyme binding site	
44004 4386 Ann Ann Ann Ann Ann Ann Ann Ann Ann Ann	1.0
tacactgott gggcagagt	19
+310+ 2387 +3312-13	
5-2112-14	

: :::

HOMA Sapiens	
<pre>cd20. cd23. Cyclin F ribozyme binding site</pre>	
<pre>&lt;:400 + 0387 gt potgasto tgttogagg</pre>	19
<pre>HD10 + D388 HD11 + 19 HD12 + DNA HD13 + Homo sapiens</pre>	
<pre><pre><ili3. binding="" cyclin="" f="" pre="" ribozyme="" site<=""></ili3.></pre></pre>	
<pre>&lt;400&gt; 0388 tgagtotgtt cgaggatga</pre>	19
<pre>#210 &gt; 1389 #211 &gt; 19 #212 &gt; DMA #213 &gt; Homo sapiens</pre>	
<pre>+330&gt; +333+Cyclin F ribozyme binding site</pre>	
-:400 - 2389 qaqtotqtto gaggatgag	19
<pre>&lt;210 &gt; 2390 &lt;2211 - 19 &lt;2212 - DNA &lt;2015 - Homo sapiens</pre>	
-Class Cyclin F ribozyme binding site	
<pre>+(4/0)+ 1/90 atdacetqtt tgaggagge</pre>	19
+02160 2391 +02110 14 +02110 DNA +02170 Homo sapiens	
+2.7C+ +2.33+ Cyclin F ribozyme binding site	
+1400 > 2.391	

tgaeetgttt gaggagget	19
<pre>&lt;210 &gt; 2392</pre> <pre>&lt;211 &gt; 19</pre>	
HODIA Homo sapiens	
*:320 ×	
RABB + Oyolin F ribozyme binding site	
-:400 + 1392	1.0
gaggotgoto atcagggat	19
H210 + 2393	
S211 + 19	
KC1C× DNA KC13> Homo sapiens	
- NETO- Times Suprems	
<220×	
<223> Cyclin F ribozyme binding site	
K400× 1393	
gotgotoato agggatgto	19
HC108	
FILLE THA	
- 113 - Homo sapiens	
-MMCB- Dyalin F ribozyme binding site	
+:400>-2394	
cagggatgto tgaccaget	19
H:010N 0395	
-0110-13-35 -0110-13-	
HOLLO DNA	
-CC138 Homo sapiens	
H226:-	
<pre>%223% Cyclin F ribozyme binding site</pre>	
and the state of t	
+:400:- 2395	
tgandagete etaceteet	19
+0:10:++36	
<pre><!--!!110</pre--></pre>	
+:21.25 DNA	
-:213: Homo sapiens	

<pre><i200> <ilu3> Tyclin F ribozyme binding site</ilu3></i200></pre>	
-0400 + 0396 -coagetesta cetestetg	19
<pre>%210 * 2397 %211 * 19 %212 * DNA %213 * Homo sapiens</pre>	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>0400 + 2397 otootacete ototgggaa</pre>	19
<pre>#210 &gt; 2398 #211 * 19 #212 &gt; DNA #213 + Homo sapiens</pre>	
+220> +223> Cyclin F ribozyme binding site	
+400 × 2398 ctacctoctc tgggaaago	19
+210 × 2399 +211 + 19 +212 × DNA +213 > Homo sapiens	
<pre>&lt;2008 </pre>	
<pre>0400+ 0399 cagatgtgtc agatcctgg</pre>	19
HI108 2400 HI119 19 HIII DNA HIII38 Homo sapiens	
HCCCOS HCCCOS Cyclin F ribozyme binding site	
-:400:- :400 gtgtcaqatc ctgggcgat	19
701 At 11101	

H211 + 19 H212 + DMA	
+213 + Homo sapiens	
+:22m3	
+3230 Gydlin F ribozyme binding site	
s(4)(0) + 2401	
requirements cacagette	19
H210 + 2402	
4211 + 19	
H21.7 > DNA	
<pre>%213 / Homo sapiens</pre>	
K220×	
<pre>&lt;223 * Cyclin F ribozyme binding site</pre>	
H400 × 2402	
tocapagett cogaaaast	19
+1.10 + 0.403	
<pre><pre><pre><pre></pre></pre></pre></pre>	
AND WILLY	
+013 - Homo sapiens	
<0.00 \$ 10.00 \$	
<pre>+1.13 * Cyclin F ribozyme kinding site</pre>	
+:400 + 2403	
ocadagetto egaaaacte	19
k(210) - 2404	
<211): 19	
<pre><c1d> INA</c1d></pre>	
<pre>si213/s Homo sapiens</pre>	
+:200;-	
<2235 Cyclin F ribozyme binding site	
\$14,000 - 24,04	
coquadaeto agggactac	19
+01103 0.405	
-0.0118-19	
-0.1110 - 0.110A	
+21.8 Homo sapiens	
+(2)10)+	
<2230 Cyclin F ribozyme binding site	

-400-2405 tolggglota ogotogoaa	19
+010+0406 <011+19 +010+0NA +010+Homo sapiens	
<pre>%200 * %203 * Cyclin F ribozyme binding site</pre>	
H400 + 0406 gastacqsts gcaaaggst	19
+010 + 2407 <0112	
<pre>#210# #223 * Cyclin F ribozyme binding site</pre>	
-0400 - 2407 ogdagotigto tittagodaa	19
+D10% D408 <d11% 19<br="">+D10 + DNA +D13 + Homo sapiens</d11%>	
<pre><pre><i220> </i220></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre> <pre></pre> <pre></pre></pre></pre>	
84008 0408 Gagotgtott tagodaaag	19
<pre>%Lius 0409 %Diis 19 %Diis ENA %Diis Homo sapiens</pre>	
<pre><pre>&lt;2200:</pre> <pre>&lt;223:- Cyclin F ribozyme binding site</pre></pre>	
-:400-::409 Agetatatt agecaaage	19
+0010+ .410 +0010+ 19 +0012+ DNA +0213> Homo sapiens	

0220 - 0223 - Oyolin F ribozyme binding site	
R400+ 2410 Watutofita godaaagad	19
0010	
<pre>#220&gt; #223&gt; Cyclin F ribozyme binding site</pre>	
<pre>-:400&gt; 2411 aaaccagctt ggactggag</pre>	19
+010 > 0412 +011 > 19 +010 > DNA +013 + Homo sapiens	
<pre>&lt;220 &gt; &lt;223   Cyclin F ribozyme binding site</pre>	
H400 - 2412 qtqaqaqott ocagtgaga	19
<pre>%010 + 2413 &lt;0211 * 19 %212 * DNA %213 * Homo sapiens</pre>	
<pre>*:220* *:223 - Tyclin F ribozyme binding site</pre>	
<pre>%400% 2413 **gagagette cagtgagat</pre>	19
+02100+ 0414 +02110+ 19 +02100+ 0NA +02130+ Homo sapiens	
+0.000+ +0.00 Cyclin F ribozyme binding site	
<pre>&lt;4000 2414 cagtgagatc gtctgccag</pre>	:9

:

<pre>&lt;(01)</pre>	
<pre>%1200 %1130</pre>	
-:400% 2415 tgagatoqto tgodagota	19
+010 + 0416 +0011 + 19 +0010 + DNA +0013 + Homo sapiens	
<pre><dii0></dii0></pre> <pre><dii3> Cyclin F ribozyme binding site</dii3></pre>	
<400> 2416 otgodagota titicaggot	19
+210 > 2417 +211 > 19 +212 > DNA +213 > Homo sapiens	
+220+ +223> Cyclin F ribozyme binding site	
-400 · 2417 godagotatt toaggotto	19
+010+ 0418 +011+ 19 +010+ TNA +010+ Homo sapiens	
k22008 k2220 Cyclin F ribozyme binding site	
-:4005-3418 ocaqotatti caggottoc	19
+00105 041+ +00105 1* +00105 LNA +00105 Homo sapiens	
-022() -0225> Cyclin F ribozyme binding site	

H400+ R419 Bag staftts aggottoco	19
H210 + 1420 H211 + 19 H212 + DNA H213 + Homo sapiens	
<pre>02209 02139 Cyclin F ribozyme binding site</pre>	
<pre>&lt;400 + 1420 ttteagqett cocaggetg</pre>	19
<pre>&lt;210 &gt; 0401 &lt;0011 &gt; 19 &lt;0010 &gt; DNA &lt;0013 &gt; Homo sapiens</pre>	
<pre><pre><pre><pre></pre> <pre><pre></pre> <pre><pre></pre> <pre>Color</pre> <pre><pre>Cyclin F ribozyme binding site</pre></pre></pre></pre></pre></pre></pre>	
<pre>+:400 * 2421 ttpaggette coaggetgt</pre>	19
<pre>K210% 0422 K211% 19 K212% DNA K213% Homo sapiens</pre>	
K2200 K2235 Cyclin F ribozyme binding site	
<400% 0400 ccaqqctqtc agtaaacaa	19
+0010 × 0.423 +0211 × 19 +0212 × DNA +0213 × Homo sapiens	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
H400H 0413 gotyteadta aacaacaag	19
+0110:+ 0424 +0.110:- 1 * +0212:+ DNA	

<pre><!--!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!</th--><th></th></pre>	
-:220 ←	
<pre><pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre>&lt;</pre></pre>	
<400 ≥ 2424	
abaabaaqtb ttotoogtg	19
+:210 + 1425	
H211 + 19	
HIDID + DNA	
<pre><!--li--><pre></pre></pre> <pre></pre>	
K220 +	
<pre></pre>	
<400% 2425	
aadaagtett otoogtgoa	19
<210> 2426	
42118 19	
KC12> DNA	
<pre><pre><pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><td></td></pre></pre></pre>	
<220 S	
<pre>&lt;223 &gt; Cyclin F ribozyme binding site</pre>	
·.400 + 0406	
abaagtette teegtgeag	19
k211> 19	
KO13 NIDNA	
<pre>&lt;013&gt; Homo sapiens</pre>	
+(2.20 %	
<pre>&lt;223 &gt; Cyclin F ribozyme binding site</pre>	
-:400.5-2427	
aaqtottoto ogtgoagaa	19
+C1.00+ D428	
(211) 19	
HILIDH DNA	
<pre><cl30 homo="" pre="" sapiens<=""></cl30></pre>	
$\cdot :: : \subseteq \Omega : \cdot$	
H2000 Cyclin F ribozyme binding site	
-:400°× 2428	
gaagggacto aatgacaca	19

+0210 + 2429 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
+1126 + +1223 + Typlin F ribozyme binding site	
-:400 + 2429 caatgaggta cattotgat	19
+:210 + 1430 +:211 + 19 +:212 + 5NA +:213 + Homo sapiens	
<220> <223> Cyclin F ribozyme binding site	
-:400:- 2430 qaqqtacatt ctgatcgac	19
<pre>&lt;210 &gt; 2431 \$211 &gt; 19 \$212 &gt; DNA \$213 &gt; Homo sapiens</pre>	
-0.20- -0.23- Cyclin F ribozyme binding site	
<pre>&lt;400 \ 0.431 aggtacatto tgatogact</pre>	19
<pre>%210</pre>	
-1110s -11110s Cyclin F ribozyme binding site	
+:400> 1432 cathotoato gaotggotg	19
+0100 0433 +0110 19 +0100 DMA +0130 Homo sapiens	
-:220:>	

	<pre><!--233   Cyclin F ribozyme binding site</pre--></pre>	
	(4.00 ± 0.433)	
	autogaaatt godaccatg	1 9
	+(210 + 3434	
	-211 · 14	
	<pre>&lt;215 - Homo sapiens</pre>	
	+200 +	
	-223 - Cyclin F ribozyme binding site	
	tyaatgactt cacaageet	1 3
	K210 + 2435	
	K111 + 19	
	KO10> DNA	
-	<pre>&lt;213&gt; Homo sapiens</pre>	
	< 3.20 >	
	<pre><pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre>&lt;</pre></pre>	
	+4005-2435	
	gaatgaette acaageetg	19
	₹2100 - 0436	
	+2115-19	
	<pre><b128 dna<="" pre=""></b128></pre>	
	<pre>%3138 Homo sapiens</pre>	
	<0.000	
	<20038 Oyolin F ribozyme binding site	
	k 4 0 00%   2 4 3 6	
	tgqaccaqta cctgcggag	19
	k(2108-)(2437	
	-02118 19	
	HINTER DIA	
	+00130 Homo sapiens	
	kinnón-	
	<pre>00.030 Cyclin F ribozyme binding site</pre>	
	+.400:+.1.437	
	tgecgeggta caggeteca	19
	KD100- 2439	
	H2119-19	

HO13: DMA HOMO sapiens	
-0020	
-(400 + 2438 gtalaaggete cagetgetg	19
+010 + 2439 +011 + 19 +0110 + DNA +013 + Homo sapiens	
-0.000 / -0.	
H4005 2439 retaggeate gootgeatg	19
+0010+ 0440 +0011> 19 +0010+ DNA +0013> Homo sapiens	
<pre>&lt;0.00 + &lt;0.03 - Cyclin F ribozyme binding site</pre>	
k:400 + 2440 Stq:atgqtc atotgcasc	19
<pre>&lt;210 - 0441 &lt;211 - 19 &lt;212 &gt; DMA &lt;213 - Homo sapiens</pre>	
-Minor- -Minor-Cyclin F ribozyme binding site	
<pre>040000 0441 0atqqteate tgcaccegg</pre>	19
+02108 [442 +0.118 10 +0.100 DNA +0.130 Homo sapiens	
<pre><dino- <223="" binding="" cyclin="" f="" pre="" ribozyme="" site<=""></dino-></pre>	
<400 - 24/12	

1 1

gbabboqgtt tatbagtaa	19
H210 + 2443 H211 + 19 H212 + DNA H213 + Homo sapiens	
00200 0003 - Oyolin F ribozyme binding site	
0400 × 2443 Gaseggttt atcagtaaa	19
+2210 > 2444 +2211 > 19 +2212 > DNA +2213 > Homo sapiens	
<220 > <223 > Cyclin F ribozyme binding site	
6400 - 2444 abboggitta itoagiaaag	19
<pre>&lt;210 + 2445 &lt;211 + 19 &lt;2212 + DNA &lt;2213 + Homo sapiens</pre>	
<pre><c20 -="" <c203="" binding="" cyclin="" f="" pre="" ribozyme="" site<=""></c20></pre>	
-:400+ :2445 coqqtttatc agtaaagag	19
+0110.+ 0446 +0111- 19 +0112- DMA +0113- Homo sapiens	
-MINOS- -MINOS- Cyclin F ribozyme binding site	
::400:- 2446 titatcaαta aagagatee	19
<pre><c100 ::19="" ::447="" ::dna="" <c110="" <c213="" <c:110=""> Homo sapiens</c100></pre>	

<pre>:(220 + :(2.03 + Cyclin F ribozyme binding site</pre>	
H400 + 2447 taaaqaqato otgaccato	19
H210 + 2448 H211 + 19	
HOMA Homo sapiens	
+220 + +223 + Dyolin F ribozyme binding site	
<pre>-:400 + 2448 cotgaccate ogggaggcc</pre>	19
<pre>&lt;310: 3449 &lt;3211:19 &lt;3212: DNA &lt;313: Homo sapiens</pre>	
<pre>#2200# #223 * Cyclin F ribozyme binding site</pre>	
-:400x ::449 agaggongta tggotoacg	19
+1210 + 2450 +1211 + 19 +1212	
+:2.20%	
<pre><pre><pre><pre></pre> <pre></pre> <pr< td=""><td></td></pr<></pre></pre></pre>	
+:400 + 2450 ogtatggoto acggacaac	19
+010+0451 +0010+03 +0010+00A	
+0130 Home sapiens	
+MLCO+ +MLCO+ Cyclin F ribozyme binding site	
-:400:- 2451 qacaacatti acaagtacg	19
+210× 2452	

+0011 · 13 +0012 · DNA +0013 · Homo sapiens	
<pre>&lt;220 * &lt;223 * Dyclin F ribozyme binding site</pre>	
<pre>0400 + 0452 abaababtta caagtacga</pre>	19
<pre>&lt;010 + 2453 &lt;011 + 19 &lt;012 + DNA +0013 * Homo sapiens</pre>	
<pre><dig0 -="" <dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"></dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor="final-right"><dicolor< td=""><td></td></dicolor<></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dicolor="final-right"></dig0></pre>	
<400> 2453 ottacaaqta ogaggacot	19
<pre>&lt;310&gt; 2454 &lt;311&gt; 19 &lt;312&gt; DNA &lt;313&gt; Homo sapiens</pre>	
<pre>&lt;120&gt; &lt;123 Gyolin F ribozyme binding site</pre>	
ा4008 2454 वृद्धपञ्चनवर्षक gtotoogoo	19
<pre>&lt;210 \ 2456 &lt;2211 \ 19 +2212 + DMA &lt;213 \ Home sapiens</pre>	
<pre><duo> still 3 - Cyclin F ribozyme binding site</duo></pre>	
<pre>&lt;400 &gt; 0455 ogagatogto teegeettg</pre>	19
+D10. D456 +D11: 19 +D1D: DNA +D1D: Homo sapiens	
+220> +223> Cyclin F ribozyme binding site	

<pre>&lt;400 &gt; 2456 agategoete egeettgga</pre>	19
H210 + 2457 H211 + 19 H212 + DNA H213 * Homo sapiens	
<pre>00.00 / 0</pre>	
<pre>0400 + 1457 teteogeout ggaagggaa</pre>	19
<pre>&lt;210&gt; 2458 &lt;2211&gt; 19 &lt;212&gt; DNA &lt;2213&gt; Homo sapiens</pre>	
<pre><d200> &lt;2223&gt; Cyclin F ribozyme binding site</d200></pre>	
स्वेग्री २ २४५८ व्यवस्थित व्यवस्थात व्यस्थात व्यवस्थात	19
+00100+ 2459 +0011+ 19 +0010> DNA +0013+ Homo sapiens	
<pre><pre><pre><pre>&lt;2220 &gt;</pre> <pre>&lt;2223 &gt; Cyclin F ribozyme binding site</pre></pre></pre></pre>	
-14000- 2459 प्रवृद्धagatto gagtococa	19
+00100+ 0460 +00110+ 19 +00100+ DNA +00130+ Homo sapiens	
<pre>&lt;2200:- &lt;2230: Cyclin F ribozyme binding site</pre>	
+:400:- 2460 gattogagto cocactgtg	19
+0.100+ 0.461 +0.110+ 0.4 +0.110+ DNA +0.213> Hemo sapiens	

<pre>&lt;2200 &lt;2223 - Cyplin F ribozyme binding site</pre>	
H400W 0461	1.0
ataatgaatt acaaggagg	19
R210 × 0462	
H211 × 19	
H212 > DNA	
CRIB+ Homo sapiens	
Hain H	
<pre><pre><pre>&lt;213 * Gyclin F ribozyme binding site</pre></pre></pre>	
<400 > 0.462	
tggtagatta caaggaggt	19
<pre>&lt;210 + 2463 &lt;&lt;211 \ 19</pre>	
<pre>&lt;211 * 19 &lt;212 * DNA</pre>	
K213 Homo sapiens	
<2200 A	
<pre><pre><pre></pre> <pre>CDD30 Cyclin F ribozyme binding site</pre></pre></pre>	
+400 × 0463	
caaqqaqqto otgotgaog	19
+0010> 0464 +0011> 19	
<010> DNA	
<pre>&lt;013&gt; Homo sapiens</pre>	
W2208	
<pre>%L23* Cyclin F ribozyme binding site</pre>	
-:400:- :2464	1.0
gatgacgata gtocotgtg	19
+122.27+ 124.65	
-02110-19	
HILLS INA	
+213+ Homo sapiens	
+(224)-	
<pre><pre><pre></pre> <pre><pre></pre> <pre>Cyclin F ribozyme binding site</pre></pre></pre></pre>	
(10%)	
-:400:- 24%5 gaogotagto cotgtggag	19
gaogocagee coegeggag	1.0

<pre>0010 - 2466 -0011 - 19 -0012 - DNA -0013 - Homo sapiens</pre>	
+0.000+ +0.000 + Gyplin F ribozyme binding site	
H400 - 2466 tytgdagath octotgdga	19
+1210 + 1467 +1211 + 19 +1213 > Bome sapiens	
<pre>&lt;220&gt; &lt;223&gt; Cyclin F ribozyme binding site</pre>	
<pre>#400+ 0467 gtgmagette ototgegag</pre>	19
<pre>&lt;210 &gt; 2468 &lt;211 \ 19 </pre> <pre>&lt;212 &gt; DNA &lt;213 &gt; Homo sapiens</pre>	
+2205 +223+ Cyclin F ribozyme binding site	
-:400.8 2468 cagettests tgcgagets	19
+0108 0469 +0110 19 +0110 DMA +0130 Homo sapiens	
22%- -:21%- Cyclin F ribozyme binding site	
<pre><id(non-24e) focotgotg<="" ofacquagetc="" pre=""></id(non-24e)></pre>	19
<pre>##1108 0470 ##1108 14 ##1108 LNA ##1130 Hemo sapiens</pre>	
<pre><pre><pre><pre></pre> <pre><pre></pre> <pre>Cyclin F ribozyme binding site</pre></pre></pre></pre></pre>	

0410 + 2470 Asgagoreta cotgotosa	19
+010 + 0471 +011 + 19 +010 + DNA	
-M13 · Homo sapiens	
+:2200 +:223 + Cyclin F ribbzyme binding site	
-:400 + 2471 obagoctato egectaege	19
<pre></pre>	
<pre>&lt;211&gt; 19</pre>	
<212> DNA <213> Homo sapiens	
<020%	
<pre><pre><pre></pre> <pre>Cubozyme binding site</pre></pre></pre>	
<:400 × 2472	
tytoogoota ogooocayo	19
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
RATA DNA	
<pre>&lt;:13&gt; Home sapiens</pre>	
<020>	
<pre><pre><d23> Cyclin F ribozyme binding site</d23></pre></pre>	
-(400» 2473	1.6
agenetgete etggecaga	19
+1 100× 2474	
(C115-14	
KILIDS DNA	
+Clib Home sapiens	
(2.20)	
<pre><pre><pre><pre>Class Cyclin F ribozyme binding site</pre></pre></pre></pre>	
H400H 2474	
tagadoadte agotgtggg	19
-:210:- 1475	
(2.11): 19	
HZ12> ENA	

::213 · Homo sapiens	
+:220 <b>-</b>	
+223 + Cyclin F ribozyme binding site	
-:400 + :3475	
gt aggazata accggatto	19
+0.10 + 2476	
±0.11 ± 19	
HILLS DNA	
+11.13 + Homo sapiens	
+11.10 ×	
<pre></pre> <pre><pre></pre> <pre>Cyclin F ribozyme binding site</pre></pre>	
-(400 > 2476)	
teadogqatt otootatga	19
esassiggate occoracya	1.5
<210> 2477	
-(2118-19)	
<212> DMA	
<213> Homo sapiens	
<3208	
<pre>&lt;223&gt; Cyclin F ribozyme binding site</pre>	
<400 → 2477	
papaggatto tootatgaa	19
K2105 2478	
<211> 19	
KOLON DIA	
<pre></pre> <pre>#2135 Homo sapiens</pre>	
<pre>&lt;113 &gt; Cyclin F ribozyme binding site</pre>	
-:4008-2478	
deggattete etatgaaga	19
-00.100+ 047#	
400.00	
HILLION DNA	
+213 - Homo sapiens	
·(200)	
-2233 Cyclin F ribozyme binding site	
<400> 2473	
gatteteeta tgaagadet	19

<pre>00:10 &gt; 2480 00:11 + 19 00:12 + DNA 00:13 + Homo sapiens</pre>	
-0020	
-:400 + 2480 tgaagasets attocotge	19
<pre>0010 + 0481 0011 + 19 0012 + DNA 0013 * Home sapiens</pre>	
<pre>0220 0223 * Cyclin F ribozyme binding site</pre>	
-:400:- 2481 agadeteatt deetgegte	19
+210 + 2482 +2211	
+0220 × +0223 + Cyclin F ribosyme binding site	
<pre>&lt;4005 0480 gadotcatto cotgogtot</pre>	19
+02100+ 2483 +02110- 19 +02120+ 00A +00130- Home sapiens	
+0000+ +0003+ Cyclin F ribozyme binding site	
+04000- 2483 topo:gogto ttgagooto	19
+0100+ 1484 +0110+ 10 +0110+ DNA +0130+ Homo sapiens	
+12202	

<pre></pre> <pre></pre> <pre>C2:23 · Cyclin F ribozyme binding site</pre>	
H400 + 2484 potgognosti gagoctoca	19
H210 + 7485 H211 + 19 H212 + DNA H213 + Hamo sapiens	
+220 + +223 + Oyclin F ribozyme binding site	
<pre>8400 + 2485 ottgagosto cataagaag</pre>	19
<pre>*210&gt; 0486  &lt;211 &gt; 19  &lt;212 + DNA  &lt;213 + Homo sapiens</pre>	
<2200 <2230 Cyclin F ribozyme binding site	
<400% 2486 agootopata agaagtgot	19
<pre>%210 % 2487 %311 % 19 %210 % DNA %213 % Homo sapiens</pre>	
<pre>+120 + +123 + Cyclin F ribozyme binding site</pre>	
+400 + 2487 adaagtgott ocatgatga	19
+ 0108 .488 + 0110 10 \$ 0110 DNA + 0130 Homo sapiens	
+0100+ +0130+ Cyclin F ribozyme binding site	
+400% 1489 gaagtgotto catgatgac	19
+E10+ 2489 +E11> 19	

.

	1212 · DMA 1213 · Homo sapiens	
	MADON MABON Cyclin F ribozyme binding site	
	:400 × 248) bbaaqgabta caggcaagt	19
	(210 = 2490) (211 = 19) (212 = DNA) (213 = Homo sapiens)	
	:220> :223> Cyclin F ribozyme binding site	
	(400> 2490 paggpaagto tototgado	19
•.	0210> 0491 0211> 19 0212> DNA 0213> Homo sapiens	
	:1208 :223    Cyclin F ribozyme binding site	
	:400% 2491 igoaagtoto totgacogo	19
•	0210> 2492 0211> 19 0012> DNA 0013> Homo sapiens	
	MACS MACS Cyclin F ribozyme binding site	
	(400× 2492 Haagtototo tgacogoog	19
•	12188 2498 1211- 19 12125 DNA 12135 Homo sapiens	
	obros. 1213: Cyalin F ribozyme binding site	
• (	400×2493	

aqcagcggtt tgaggacaa	19
+1310 + 2494	
4221 - 19	
S012 > DMA	
+1213 + Homo sapiens	
+13.20 +	
-AAAA - Cyclin F ribozyme binding site	
with a problem and the problem of th	
<14:10 + 14:94	
geagegattt gaggacaag	19
K210 + 3495	
K211 × 19	
K212× DNA	
<213 Homo sapiens	
•	
<220>	
<pre>CDUB&gt; Cyclin F ribozyme binding site</pre>	
<400 × 2495	
acaagogota tggagaaat	19
<210 + 2496	
<211 > 19	
SINIA DNA	
Addiso Homo sapiens	
+1.0.1(0)+	
k400% 0496	
tgyagaaato agocaggaa	19
si210+ 2497	
**E10** 19**	
KU 103 DDA	
+111130 Hemo sapiens	
+:2.20:-	
+:223: Cyclin F ribozyme binding site	
-:4005-:2497	
tgcfgagcta cagccagtt	19
KIII 101: 1498	
*COLD:= 19	
H212H DNA	
4213> Homo sapiens	

<pre>&lt;220. &lt;222. Oyolin F ribozyme binding site</pre>	
-(4)(0 + 2498) abagboagtt gtgtgotgo	19
KU10 + 2499 HU11 + 19 KU12 + DNA	
+213 - Homo sapiens	
REEDO	
<400> 2499 qtqctgcatt aggagtgac	19
<010 - 2500 <011 > 19	
<pre>&lt;212&gt; DNA &lt;213&gt; Homo sapiens</pre>	
-:2205 -:223> Cyclin F ribozyme binding site	
<pre>+(400 \ [500] tghtgcatta ggagtgada</pre>	19
+2108 3501 +211 - 19	
<pre>&lt;211&gt; DNA &lt;213&gt; Homo sapiens</pre>	
<pre>*ALTON Cyclin F ribozyme binding site</pre>	
H4/m- 2001 Googgaatt tectcagea	19
+(010)+ 2502 +(011)+ 14	
FULLUS DNA FULLUS Homo sapiens	
+1705 +17.30 Cyclin F ribozyme binding site	
<ul><li>4400 - 3502</li></ul>	
coccqacttt cotcagoac	19
+C210: 2503	

±211 + 19	
::212 + DNA	
-3213 · Homo sapiens	
.210, nome sabiens	
- 200	
<220 *	
+0.023 + 0yclin F ribozyme binding	site
+(400 + 1503	
occyacitto otcagoaca	19
$\pm 210 \pm 2504$	
(211 + 19)	
-:212 - DNA	
<pre>%213 * Home sapiens</pre>	
•	
K220x	
<pre>&gt;223&gt; Cyclin F ribozyme binding</pre>	site
and a series of the series of	
<400 > 2504	
	19
gaettteete ageaeaggg	13
K.010 # 2505	
<211 * 19	
KO10+ DNA	
<213 Homo sapiens	
+22.20 ·	
-00.03 Gyolin F ribozyme binding	site
<400 + 2505	
aqqqqaqato cacgootto	19
<210% 2506	
H2119 19	
<pre></pre>	
<pre><pre></pre><pre></pre><pre>#21:00 Homo sapiens</pre></pre>	
•	
K2.200	
+:223: Cyclin F ribozyme binding	site
+(400) - 2506	
todaognott octoagoto	19
tibbang.diss coucage to	1.2
HD101- 2507	
+0.10 · 19	
HILION LMA	
elle Homo sapiens	
•CD2(**	
- 1223> Cyclin F ribozyme binding	site

<pre>(400 + 2507 prapagentte eteagetht</pre>	19
+0210 + 0508 +0011 + 19 +0010 + DNA +0013 + Homo sapiens	
+C220 + +C223 + Cyclin F ribozyme binding site	
H400 - 2508 squattests agototose	19
<pre>\$210 + 2509 \$211 &gt; 19 \$212 &gt; DNA \$213 &gt; Homo sapiens</pre>	
<pre>&lt;220s &lt;223 - Cyclin F ribozyme binding site</pre>	
<400> 2509 tootbagets teecteggg	19
<pre>K210 &gt; 2510 4211 &gt; 19 K212 &gt; ENA 4213 &gt; Homo sapiens</pre>	
<pre>&lt;2200x &lt;223&gt; Cyclin F ribozyme binding site</pre>	
<pre>24008 2510 ptoAgototo cotoggggo</pre>	19
<pre>%210 % 2511 %211 % 19 %212 % DMA %213 % Homo sapiens</pre>	
- AMED: -AMED: -	
<pre>+400+ L511 gototocete ggggcggag</pre>	19
+00100+0512 +0010+19 +0010+ DMA +0013+ Homo sapiens	

+:220 · +:223 · Cyclin F ribozyme binding site	
-0400 + 2512 gaadagbato caggaagac	19
	1 -/
H210 + 2513 H221 + 19	
HOUSE DNA	
+CC13 + Homo sapiens	
-titili) -	
+CLS+ Cyclin F ribozyme binding site	
<400> 2513	1.0
daggbagett egttaceae	19
K210 8 2514 K311 > 19	
HOLOS DNA	
+2213 Homo sapiens	
K000+	
<pre><!--:C23--> Cyclin F ribezyme binding site</pre>	
-0400 + 0514	
addeagette gttaccace	19
-1710H 1515	
-X112 19 -X112> DNA	
<pre>S213&gt; Romo sapiens</pre>	
+1220th	
+CCSS Cyclin F ribozyme binding site	
-:400> 1515	
dagettegtt accadeded	19
HODIOH 2516	
HOLLOW 19 HOLLOW DNA	
HC1130 Homo sapiens	
-:::::::::::::::::::::::::::::::::::::	
-CMM30- Cyclin F ribozyme binding site	
+:400:2516	
agottogtta ccaccecca	19

+:210 + 0517 +:211 + 19	
-:212 - 5NA	
+:213 · Homo sapiens	
.30	
-:220 -	
-1223 - Cyclin F ribozyme binding site	
<400 + 3517	
oggadotyto cagodagga	19
FC[10 + 2518]	
₹Д11 × 19	
HOID A DWA	
<pre>H2132 Homo sapiens</pre>	
<2105	
<pre>&lt;123&gt; Cyclin F ribozyme binding site</pre>	
<4400 · 2518	
taggacagett cotogactg	19
<210 x 0519	
<0.11x 19	
KOLDS DNA	
<pre>&lt;013 - Homo sapiens</pre>	
•	
K220 /	
-1223 - Cyclin F ribozyme binding site	
-:400 + 2519	
gggbagette etegactgg	19
H210H 2520	
(211): 19	
HOILH DNA	
<pre>+22130 Homo sapiens</pre>	
•	
+12200+	
·C23: Cyclin F ribozyme binding site	
, , , , , , , , , , , , , , , , , , , ,	
+:400+ 2520	
dagottocto gactggago	19
· · · · · · · · · · · · · · · · · · ·	
(216) 7521	
*III: 19	
MILD DNA	
:113: Homo sapiens	
·.2200·	
<pre>4223&gt; Cyclin F ribozyme binding site</pre>	

-:460> 2521	
actgotypto tggotatga	19
€210 + 2522	
<:211 • 13	
H212 > DNA	
+213 + Homo sapiens	
< 2.200 -	
<pre>*Ad3 - Cyclin F ribozyme binding site</pre>	
×400 + 2522	
gototggota tgaaggoga	19
<010 + 2523	
₹211 + 19	
<pre>&lt;012 &gt; DNA</pre>	
<pre>&lt;113 Homo sapiens</pre>	
< 2.10 -	
<pre>&lt;2003 - Cyclin F ribozyme binding site</pre>	
<400 + 2523	
gtqadagetid odagoggda	19
4210 + 0524	
+211 + 19	
-212 - DNA	
-113 - Hemp sapiens	
21% nins suprons	
<220 ×	
<pre>&lt;323 - Gyolin F ribozyme binding site</pre>	
+ 400 + 2824	
dagangeato otogatgto	19
*210 + 2525	
-211 + 19	
1011 - TMA	
-213. Homo sapiens	
S2007-	
·203 · Cyclin F ribozyme binding site	
- 400% 2505	
	10
dagdaton n gatgtoaco	19
02100-2026	
(IIII) 14	
<212> LNA	

K213> Homo sapiens	
+12.20 +	
-MRB+ Cyclin F ribbzyme binding site	
+400+ 2526	
actical acceptage to	19
+210 + 2527	
P211 - 19	
S212 + DNA	
+213 + Homo sapiens	
+2203	
<223 · Cyclin F ribozyme binding site	
<400 × 2527	
daddqtqqtd tacotgaad	19
K210 × 2528	
+1311 + 19	
<213> DNA	
<pre>si213 &gt; Homo sapiens</pre>	
<229 ×	
<pre>SOL3 + Cyclin F ribozyme binding site</pre>	
+(400 × .7508	
poquagticta estgaacse	19
K210% 0529	
<2115-19	
SCIEN DNA	
-213 - Homo sapiens	
editos	
*AAAA Cyclin F ribozyme binding site	
440 % (2829)	
gaanagnatt gotgooagg	19
4216: 2530	
R2119-19	
FUNILIFIEDNA	
+DDLS+ Homo sapiens	
+121 (.)+	
-27.55 Cyclin F ribozyme binding site	
-:400% 2530	
gccaggaatc cagtgatga	19

<pre>+010+ 2531 +011+ 19 +011+ DNA +013+ Homo sapiens</pre>	
+220 + + + + + + + + + + + + + + + + + +	
w400 + 2531 gaggaggott gtocagagg	19
<pre><d10 +="" 0530="" 19="" 8011="" <="" <d12="" <d213="" dna="" homo="" pre="" sapiens<=""></d10></pre>	
<pre>&lt;220 + &lt;223 + Cyclin F ribozyme binding site</pre>	
र्थ00 - 2532 gaggettgte cagaggeaa	19
<210 - 2533 <211 - 19 <212 - DNA <213 - Homo sapiens	
<pre>%220 * %223 * Cyclin F ribozyme binding site</pre>	
k4005-2533 cacccagato cotgoaacc	19
<pre>&lt;2108 2534 &lt;2110 19 &lt;21.0 DNA &lt;21.3 Home sapiens</pre>	
HIDDE: HIDDE: Cyclin F ribozyme binding site	
04000-0534 accordante egeaceage	19
+0.100+ 0.535 +0.110+ 19 +0.210+ DNA +0.213+ Homo sapiens	
ezen.	

42.3 - Gyolin F ribozyme binding site	
+:400 + 2535 давидардыр acgaccupa	19
+0:10 + 0:536 +0:11 + 19 +0:10 + DNA	
Homo sapiens	
+:220 + +:223 + Cyclin F ribosyme binding site	
k400 + 2536 toacgacoto agggtacto	19
<210 × 2537 <211 × 19	
<pre>&lt;310 &gt; EMA &lt;3113 - Homo sapiens</pre>	
<pre><d20< td=""><td></td></d20<></pre>	
<400 - 2537 deteaggeta etectoogt	19
x010 + 0538 x011 + 19	
RU10 + DMA RU130 Homo sapiens	
<pre>&lt;1108 &lt;1238 Cyclin F ribezyme binding site</pre>	
<pre>&lt;4000 - 0539 baggqtabto otoogtcag</pre>	19
HITOH I539 HITOH I9 HITOH DNA	
*213 Homo sapiens	
+2100+ +2230+ Cyclin F ribozyme binding site	
-:400% 2539 gqtactocto ogtoagoac	19
::210 + 2540	
(211) 19	

+0.212 + DNA	
+1213 · Homo sapiens	
+12.20 x	
<pre><pre><pre><pre></pre></pre><pre><pre></pre></pre><pre></pre></pre><pre></pre><pre></pre></pre> <pre></pre> <pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><p< td=""><td></td></p<></pre></pre>	
<4400 > 2540	
	1.0
ntodtodato agcacogoa	19
- 31 A 31 A.1	
+2210 > 2541	
-0.115 19	
HILLS DNA	
HD139 Home sapiens	
+111.10 >	
-CAS> Cyclin F ribozyme binding site	
+:400> 2541	
adogoaayto coacaagot	19
+:210× 2542	
<211> 19	
SD1D > DNA	
<pre>&lt;0.13&gt; Homo sapiens</pre>	
AMAGE Cyplin F ribozyme binding site	
<4008 2842	
odadaaqato ogtggaogg	19
KI210>   2543	
<2115-19	
R2128 DNA	
<pre>&lt;213  Homo sapiens</pre>	
-1.12ms	
-0.035 Cyclin F ribozyme binding site	
<4000 2543	
acqqtgqctt gggggccct	19
+123.0%   25.4.4	
KII 18-19	
HIDDE DNA	
KL130 Homo sapiens	
enn Co	
HZDAN Cyclin F ribozyme binding site	
H400% 2544	

dechaaceta ecteagtge	19
00010 + 2545 00011 + 19 00012 + DNA 00013 + Homo sapiens	
<pre>HLD3+ Typlin F ribozyme binding site</pre>	
k400 - 2545 aadotadoto agtgotgto	19
+0010.4 0546 +0010.4 19 +0010.4 DNA +0010.4 Homo sapiens	
<pre><pre><pre><pre><pre></pre> <pre><pre>Company the company to t</pre></pre></pre></pre></pre></pre>	
<400% 2546 cagtgotgte cotgeacag	19
+00100+2547 <0010+19 +0010+000A +00130+Homo sapiens	
RDDD: RDDB: Cyclin F ribozyme binding site	
k:4005-2547 adaqtgadto gdadadaba	19
HL100-7548 HL100-13 HL100-CMA HL130-Homo sapiens	
<pre><pre><i200:- <pre=""></i200:-></pre> <pre></pre> <pre></pre> <pre></pre> <pre>Cyclin F ribozyme binding site</pre></pre>	
(400): U548 tigodachatic aggodagga	19
<pre>40.100 .549 40.110 14 40.111</pre>	

<pre>%200 * %203 * Type Type District Type Type District Type Type District Type Type District Type Type Type Type Type Type Type Type</pre>	
(4n)+ 2549 эра маадта atgtttaca	19
#210 + 2550 #211 + 19 #212 + DNA #213 + Homo sapiens	
HODD - HODD F riboxyme binding site	
H400 + 2550 augtoatgtt tacagtgto	19
<pre>#310 x 2551 #311 &gt; 19 #313 &gt; Homo sapiens</pre>	
<pre>8000&gt; 8000</pre>	
-0400 + 2551 agtpatyttt acagtgtog	19
+0010 + 2552 +0011 > 19 +0010 + DNA +0013 + Homo sapiens	
<pre>&lt;220 * &lt;223 * Cyclin F ribbzyme binding site</pre>	
<pre>&lt;400 2552 qtcatgttta cagtgtcgt</pre>	19
+0010 + 0053 +0010 + 19 +0010 + DNA +0013 + Homo sapiens	
<pre>*CECO:- *CECO:- *</pre>	
H400H 1553 Htanagtgto gtoccocaa	19
.: 21 m - 255 #	

-:011 + 19	
-(210 + DNA	
<pre><!--il3+ Homo sapiens</pre--></pre>	
(12).	
-D23+ Cyclin F ribozyme binding site	
<ul><li>(400 + 1954)</li></ul>	
daqtiqtoqto doccaagto	19
H210 + 2555	
K211 + 19	
HIII TO	
-213 - Homo sapiens	
Clur Ruley Sapiens	
H2U0 ×	
-333 - Gydlin F ribozyme binding site	
4	
· 400 · 2555	
peopeaagte coccggaga	1 9
K210 + 2556	
4311 × 19	
COLON DNA	
<pre>&lt;213 Homo sapiens</pre>	
41.0 · 4203 · Cyclin F ribozyme binding site	
-1203 - Cyclin F ribozyme binding site	
-1203 - Cyclin F ribozyme binding site	19
0203% Dyclin F ribozyme binding site 0400% 2556 qaqqaqtqtt occcagoaa	19
<pre>CD03+ Dyclin F ribozyme binding site  &lt;400+ 2556  qaqcagtqtt ccccagcaa </pre>	19
<pre>CD03+ Tyclin F ribozyme binding site  &lt;400+ D556  qaqcagtqtt ccccagcaa  &lt;210+ D557 &lt;211&gt; 19</pre>	19
<pre>CD03+ Dyclin F ribozyme binding site  &lt;400+ 2556  qaqcagtqtt ccccagcaa </pre>	19
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<pre>CD03+ Dyclin F ribozyme binding site  C400+ 2556  gageagtgtt ccccageaa  C210+ 2557  C211+ 19  C211+ DNA</pre>	19
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CD03+ Cyclin F ribozyme binding site  C400+ 2556  Qaqcagtqtt ccccagcaa  CD10+ 2557  CD11+ 19  CD11+ DNA  CD13- Homo sapiens	19
#2003 # Cyclin F ribozyme binding site  #400 + 2556  #agageagtatt coccagoaa  #2210 + 2557  #2211 * 19  #2212 DNA  #2213 Homo sapiens	19
CD03+ Cyclin F ribozyme binding site  C400+ 2556  Qaqcagtqtt ccccagcaa  CD10+ 2557  CD11+ 19  CD11+ DNA  CD13- Homo sapiens	19
#2003 Dyclin F ribozyme binding site  #400 - 2556  gameagtatt coccageaa  #210 - 2557  #2115 19  #2115 DNA  #2113 Homo sapiens  #2105 Cyclin F ribozyme binding site	19
#2003 Dyclin F ribozyme binding site  #400 - 2556  qaqcagtqtt ccccagcaa  #210 > 2557  #2110 19  #2110 DNA  #213 Homo sapiens  #2100 Cyclin F ribozyme binding site	
#2003 Dyclin F ribozyme binding site  #400 - 2556  #agageagtatt coccageaa  #210 - 2557  #2115 19  #2115 DNA  #2115 Home sapiens  #2105 Cyclin F ribozyme binding site  #4008 7857  #agaagtatte cocageaac	
#2003 Dyclin F ribozyme binding site  #400 - 2556  gaggagtitt coccagoaa  #210 - 2557  #211	
#2003 Dyclin F ribozyme binding site  #400 - 2556  gagcagtatt coccagcaa  #210 - 2557  #2110 DNA  #213 Homo sapiens  #210 - Cyclin F ribozyme binding site  #400 - 2557  #400 - 2558  #2110 - 2558  #2110 - 2558	
C203	
#2003 Dyclin F ribozyme binding site  #400 - 2556  gagcagtatt coccagcaa  #210 - 2557  #2110 DNA  #213 Homo sapiens  #210 - Cyclin F ribozyme binding site  #400 - 2557  #400 - 2558  #2110 - 2558  #2110 - 2558	
C203. Cyclin F ribozyme binding site  (400. 2556  gaggaggtytt coccagoaa  (210. 2557  (211. 19  (211. DNA (213. Homo sapiens (210. Cyclin F ribozyme binding site  (400. 2557  (400. 2557  (400. 2558 (211. 19 (211	
C203	

::400 + 2550	
पुत्रवाप्रदेशकार्यः aacctatge	19
(210 - 2559	
H2115 19 H212 - DNA	
-1.213 - Homo sapiens	
<220 +	
<pre>&lt;223 - Dyplin F ribozyme binding site</pre>	
-(400 + 2559	
dataaachta tgcatacac	19
<210 × 2560	
<pre>&lt;211 * 19 </pre>	
<213 · Homo sapiens	
<220 ·	
<pre><!--223 - Cyclin F ribozyme binding site</pre--></pre>	
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octatgcata cacagtgag	19
<210% 2561 <211% 19	
KOIDE DNA	
<pre>S2130 Home sapiens</pre>	
-17.1.0 -	
SLLD3 S Cyclin F riboxyme binding site	
-:400h 1561	1.0
cotqqqoott gtgaggotq	19
+:210 + 1562	
HOLLS 19 HOLLS DMA	
<pre>SC13: Home sapiens</pre>	
s.2200-	
+2230 Cyclin G1 ribozyme binding site	
+:400:- 2562	
dagaagaatt gtotoggog	19
+02100+ 1563 +02100+ 14	
-0.011.0-1.4 -0.011.0-1.00A	
+2132 Homo sapiens	

-:220 ·	
<pre><d23 +="" binding="" cyclin="" g1="" pre="" ribozyme="" site<=""></d23></pre>	
<400 ≠ 2563	
objaithate teggeggeg	19
+1.10 + 1.564	
R211 + 19	
<212 + DNA	
<pre><!--!!! Homo sapiens</pre--></pre>	
+220 +	
<223 - Cyclin G1 ribozyme binding site	
<400 + 2564	
bogaggooto cabccagga	19
osyayysses cascoayya	13/
<210× 2565	
<211> 19	
KO125 DNA	
<pre>&lt;013 / Homo sapiens</pre>	
K020%	
<pre>&lt;3238 Gyelin G1 ribozyme binding site</pre>	
-:4000- 2565	
caggacagte ecotococ	19
H310N 2566	
·211> 19	
KOTON DNA	
<pre>&lt;2130 Homo sapiens</pre>	
K000)	
+3235 Syclin G1 ribozyme binding site	
-:400% (566	
Agtoccatic coogggest	19
<22105-1567	
H2113-19	
KILIZE DNA	
C2135 Homo sapiens	
chines	
H27 Fr Cyclin G1 ribozyme binding site	
,	
<4001- 2567	
accgggeeta tatastatt	19

<pre>#210 &gt; 2568 #2211 + 19 #2212 + DNA #2213 + Homo sapiens</pre>	
<pre>&lt;220 * &lt;223 * Cyclin G1 ribozyme binding site</pre>	
्येश्वर १५६६ अववृक्ष्याच्या testettge	19
+0010 + 0569 +0011 + 19 +0010 + DNA +0013 + Homo sapiens	
<220 · <221 · Cyclin G1 ribozyme binding site	
-:400.5 2569 agaatatata atattgaat	19
<210: 2570 <211: 19 <212: DNA <213: Home sapiens	
<pre>&lt;2200: &lt;2220: &lt;2223: Cyclin G1 riboxyme binding site</pre>	
<pre>ki400:- 2570 ctctctcctc ttgcctacg</pre>	19
<pre>&lt;:210: 2571  <c110: 14="" <c120:="" <c130:="" dna="" home="" pre="" sapiens<=""></c110:></pre>	
-00000- -00030-Cyclin G1 ribozyme binding site	
-:4002571 	19
+02100-0472 +0210-14 +0210- DNA +02180- Homo sapiens	
+:220+ +:223> Cyclin G1 ribozyme binding site	

.

0400 × 2572 Stobigosta ogagtocco	19
K210 + 3573 K211 + 10 K212 + DNA	
-C13 - Homo sapiens	
+220+ +223+ Cyclin G1 ribbzyme binding site	
+400 × 1575 Gottaggagto codototoo	19
<pre> <pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>0000 complete 0000 comple</pre>	
advecedte testegtag	19
<pre>#210 - 2575 #211 - 19 #212 - DNA #213 - Homo sapiens</pre>	
<pre>-/2108 </pre> <pre></pre> <pre>Cyclin G1 ribozyme binding site</pre>	
<pre>s400b 2575 tecccctote ctogtagge</pre>	19
+01100+ 0576 +0110- 19 +0110+ ENA +0130- Homo sapiens	
-02200- -02230- Cyclin G1 ribozyme binding site	
H40H+ 2576 chatathata gtaggaata	19
HILLON 2577 HILLON 19 HILLON DNA	

.

<pre></pre>	
<pre>-mm0mm2 - Cyclin G1 ribozyme binding site</pre>	
+(400 + 2577) totostigta ggoototog	19
HILLON 0578 HILLON 18 HILLON DNA	
<pre>Homo sapiens</pre>	
<pre>%2.00 / %Ldr&gt; Cyclin G1 ribozyme binding site</pre>	
<pre>&lt;400% 2578 ogtaggosts teggatetg</pre>	19
<pre>H210&gt; 2579 H211 &gt; 19 H213 &gt; DNA K213 &gt; Hamo sapiens</pre>	
<pre>&lt;320&gt; &lt;3210</pre>	
-:400 - 2579 tangoctoto ggatotgat	19
<pre><pre><pre>&lt;0:10 + 2580 &lt;0:11 &gt; 19 &lt;0:11 &gt; DNA &lt;0:13 &gt; Homo sapiens</pre></pre></pre>	
-:226:223 - Cyclin G1 ribczyme binding site	
H400 - 1580 ototogdato tgatatogt	19
+0210 + 1581 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
+:210:- +:213:- Cyclin G1 ribozyme binding site	
+:400% 2081 ggatetgata tegtggggt	19

<pre>4010 + 2582 4011 + 19 4010 + DNA 4013 + Homo sapiens</pre>	
HU20+ HU23+ Dyclin G1 ribozyme binding site	
<pre>+(4005 0580 atotgatato gtggggtga</pre>	19
+1010 + 0583 +1011 + 19 +1011 > DNA +1013 > Homo sapiens	
+2205 +2238 Cyclin G1 ribozyme binding site	
-:400 × 2583 quqqqtqqtt accgtgagg	19
<pre><c10> 2584 <c211> 19 <c112 +="" <c213="" dna="" homo="" pre="" sapiens<=""></c112></c211></c10></pre>	
+220 + +223 + Cyclin G1 ribczyme binding site	
<400% 2584 addqtgqtta cogtgagga	19
+0100+ 2589 +0110+ 19 +0110+ DMA +0110+ Home sapiens	
<pre>+0.200+ +0.000+ Cyclin Gl ribozyme binding site</pre>	
<pre>&lt;4000 .585 agotgoagto totgtoaag</pre>	19
<pre>#1100</pre>	
+:22 <del>0</del> :>	

<pre>&lt;:223 · Cyclin 31 ribozyme binding site</pre>	
<pre>:(4:00 + .1586 stgcagtsto tgtcaagat</pre>	19
+0210 + 2537 +0211 + 13	
+C212 + DNA +C213 + Homo sapiens	
+:220 + +:223 + Cyclin G1 ribozyme binding site	
+:400 + 2587	
agtototyto aagatgata	19
+:210 + 2588 +:211 + 19	
+C113 + DNA +C113 × Homo sapiens	
+0.20 + +0.23 + Cyclin G1 ribozyme binding site	
-:4002588 -caaqatqata gaggtactg	19
H2105 2589 H2115 19	
HOLDS DNA	
+013% Homo sapiens	
+:220×	
+Clift Cyclin G1 ribozyme binding site	
H4000- U589 datadaddta otgacaaca	19
walawawyta ocyacaaca	10
KD108-0890	
(211): 19	
HC112H ENA HC13H Home sapiens	
-0.230 Cyclin G1 ribozyme binding site	
+:40g>590	
daadtgadtd toagaaact	19
₹210% 2591	
+121114 1 )	

H212 + H113 +	DNA Homo sapiens	
H220 +	Cyclin G1 ribozyme binding site	
4400. aptyac		19
-0.10 - -0.11 - -0.12 - -0.13 -	19	
H220×	Cyclin G1 ribozyme binding site	
<400> gaaact		19
<210 × <211 > <212 > <212 > <213 >	19	
H220≯ H223⊁	Oyolin G1 ribozyme binding site	
<pre></pre> <pre>atgood</pre>		19
<210> -011> -012> -012>	19	
0220s 0223s	Cyclin G1 ribozyme binding site	
-14005 aacagg		L 9
+10.105 +10.115 +10.115 +10.135	19	
·021.30	Cyclin G1 ribozyme binding site	
-14000	2595	

caggagtota gatgtcago	19
<pre>#010 + 2596 #011 + 1# #012 + DNA #0013 + Homo sapiens</pre>	
<pre>&lt;0.00 * &lt;0.03 * Gydlin G1 ribozyme binding site</pre>	
-400 - 2596 Fotagatifto agodaaagg	19
+210 + 2597 +211 + 19 +212 + DNA +213 > Homo sapiens	
<pre>&lt;220 • &lt;223 • Cyclin G1 ribozyme binding site</pre>	
k400 + 2597 qodaAaqqtd tgtggtttg	19
<pre>&lt;210 &gt; 2598 &lt;2211 &gt; 19 &lt;2212 - DNA &lt;2213 - Homo sapiens</pre>	
<pre>&lt;220 • &lt;223 • Cyclin G1 ribozyme binding site</pre>	
ार्थ00 + 2598 प्राट्यप्रप्राप्य tu tgagactaa	19
<pre>0010 + 0599 0011 + 19 0012 + DNA 0013 + Homo sapiens</pre>	
+COLOR- +COLOR-Cyclin G1 ribozyme binding site	
-:400:- 2599 totgtgdttt gagactaat	19
<pre>### 1005 1x00 ### 1105 1x ### 1215 DNA #### S2130 Homo sapiens</pre>	

<pre> character character</pre>	
<pre>0400 + 1600 uttragabta attgagtot</pre>	19
0010 + 2601 0011 + 19 0010 + DNA 0011 = Himo sapiens	
HIBBO - HIBBO - Cyclin G1 ribozyme binding site	
୍ୟ-୧୦୧୬ ଅଟିପୀ gagactaatt gagtotgoa	19
<pre>#210% 2602 &lt;221% 19 &lt;221% ENA &lt;221% Homo sapiens</pre>	
<pre><!--ldo. <!documents</th--><th></th></pre>	
K400+ 1602 taattgadto tgoadadga	19
<pre>&lt;0.10 + 0.603 +0.11 + 19 +0.12 + DNA &lt;0.2130 Homo sapiens</pre>	
k0100+ k013+ Gyolin G1 ribozyme binding site	
-:400:- 1603 gnanacquia atggeetea	19
\$1210 + 1404 \$1211 + 19 \$1212 + DNA \$1213 + Homo sapiens	
SMM 000 SMMSO Cyclin G1 ribozyme binding site	
-:400 - 2004 taatogooto agaatgact	19
+:210> 2605	

H211 + 19 H212 + DMA H213 + Homo sapiens	
<pre>0220 - 0203 - Gyolin G1 ribozyme binding site</pre>	
୍ୟପତି - ଅଟ୍ଡିଡି tgoaagacta agggacttt	19
+0.10 + 0.606 +0.11 + 19 +0.12 + DNA +0.13 + Homo sapiens	
<pre><d200x <d223="" binding="" g1="" oyclin="" pre="" ribozyme="" site<=""></d200x></pre>	
-:400> 2606 taagggaott tgaagtaaa	19
+310 > 1607 +3212 > 19 +3212 > DNA +3213 > Homo sapiens	
Riggos Riggos Cyclin G1 ribozyme binding site	
+400> 1607 sagggaettt gaagtaaaa	19
<pre>%0108 2608 %0115 14 %0110   TNA %0113   Homo sapiens</pre>	
+:2205 +:223+ Cyclin G1 ribozyme binding site	
ក1400% 1.608 cfittqaaqta aaagatott	19
<pre>%2108 1 009 %2118 1 0 %2120 DNA %2120 Home sapiens</pre>	
+22(C)+ +22(B)> Cyclin G1 ribozyme binding site	

H400% 2609 qtaawaqato ttottagto	19
+0210 + 0610 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
+MARY+ +MARY+ Cyclin G1 ribozyme binding site	
0400 - 1610 asasyatuti ottagtota	19
<pre><c10 +="" 0611="" <c11="19" <c12="DNA" <c13="" =="" homo="" pre="" sapiens<=""></c10></pre>	
<pre>&lt;220 * sub3 * Cyclin G1 ribozyme binding site</pre>	
<400% 3611 aaagatotto ttagtotaa	19
<pre>&lt;210</pre>	
+1213 + Home sapiens	
<pre>00.00 + 00.03</pre>	
<pre>44008 0610 agatotfott agtotaact</pre>	19
0.108 2613 00118 19 00118 DNA 0218 Homo sapiens	
-0100- -0203- Cyclin G1 ribozyme binding site	
44000- 2x15	
gatominita giotaacio	19
+0.100+ [e14 +0.110+ [+ +0.120+ DNA +0.130+ Homo sapiens	

<2200 <2203 - Cyclin G1 ribozyme binding site	
H400 + 2614 SttStt9gts taactcagt	19
<pre><pre><pre><pre><pre></pre></pre></pre></pre></pre>	
+0.11. + DNA +0.13 + Homo sapiens	
-ummon -ummon Gyolin G1 ribozyme binding site	
+:400> 2615 tottagtota actoagtto	19
<pre>&lt;210&gt; 2616 &lt;211&gt; 19</pre>	
<pre>x212 + DNA x213 + Homo sapiens</pre>	
<pre>%220% %LL3% Cyclin G1 ribozyme binding site</pre>	
<400 \ 1616 agtotaacto agttotttg	19
+:210+ 2617 +:211 + 19	
S212> DNA <2138 Homo sapiens	
<pre>+ihhe - +ihhe - yolin G1 ribozyme binding site</pre>	
+:400% 2617 taactcagtt ctttggctt	19
+07105 7618 +07105 14	
+D 13% Homo sapiens	
+:110:- +:113 · Cyclin G1 ribozyme binding site	
1400 > 2618 aactcagtto tttggcttt	19

<pre>4210 - 2619 4211 - 14 4212 - DNA 4213 - Homo sapiens</pre>	
-0.230 ·	
-:400 + 2619 otoaqttott tggotttga	19
<pre>&lt;210 &gt; 0620 &lt;211 + 19 &lt;210 + DNA &lt;213 &gt; Homo sapiens</pre>	
<2005 <2005 Cyclin G1 ribozyme binding site	
k400% 2620 toaqttottt ggotttgac	19
<pre>&lt;210% 2621 &lt;211% 19 &lt;212% DNA &lt;213% Homo sapiens</pre>	
<pre>&lt;210. &lt;2230 Cyclin G1 ribozyme binding site</pre>	
k:400> 2021 tetttggett tgacacaga	19
+0110+ 0602 +0110+ 19 +0110+ DNA *013+ Homo sapiens	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>+14000+ 2672 offfdgetit gaeacagag</pre>	19
+00100 1623 +00100 14 +00100 DNA +00130 Homo sapiens	
+1.100+ +11230 Cyclin G1 ribozyme binding site	

-(4))) - 1623 bagayasatt ttototago	19
<pre>&lt;(210 + 2624 &lt;(211 + 19 &lt;(212 + DNA &lt;(213 + Homo sapiens</pre>	
২০০০ ২০০৪ - Cyclin G1 ribozyme binding site	
k(400) 2024 agagacattt tototagot	19
<210 × 1625 <211 × 19 <212 + DNA <213 * Homo sapiens	
<2000 s <2003 · Cyclin G1 ribozyme binding site	
<400% 2625 gagacatttt ototagotg	19
<pre>&lt;210</pre>	
<pre>&lt;2000 &lt;0.030 Cyclin G1 ribozyme binding site</pre>	
R4008 2616 agadatitic totagotgt	19
<pre> <pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	
SUMMODE Cyclin G1 ribozyme binding site	
Higure 2627 Adarkticic tagotigtga	19
+00100+ 1608 +02100+ 19 +0212> DNA	

<pre>&lt;213&gt; Homo sapiens</pre>	
+:220:+	
-1233 - Cyclin Gl ribozyme binding site	
+14000+ 2628	
arttictota gotgtgaat	19
HIL10H IMI9	
-2115-19	
-11.12× 5NA	
-C13× Homo sapiens	
+220×	
<pre>-223&gt; Cyclin G1 ribozyme binding site</pre>	
-(400x J679	
gotgtgaatt tactggaca	19
+010× 1630	
×211> 19	
<2125 DNA	
+013> Homo sapiens	
- 1200×	
-2035 Cyclin G1 ribozyme binding site	
+ 400% 2630	
etgtgaatit aetggaeag	19
k216N-2631	
K211N 19	
KIDION TNA	
+C13> Homo sapiens	
+D208	
+CDBS Cyclin G1 ribozyme binding site	
-:400:- 1631	
tgtgaattta otggacaga	19
x(210)5 (26.32)	
+2118-19	
ROIES UNA	
+1.13+ Homo sapiens	
KUIZ IA K	
<pre></pre> <pre>&lt;2.:3 · Cyclin Gl ribozyme binding site</pre>	
tggacagatt cotgtotaa	19

<pre>%210 * 0633 %211 * 13 %212 * DNA %213 * Homo sapiens</pre>	
+:220 + +:: -::::::::::::::::::::::::::::::::	
0400 + In33 ggadagatto otgtotaaa	19
<pre>&lt;:210 + 3634 &lt;:211 + 19 &lt;:212 &gt; DNA &lt;:213 &gt; Homo sapiens</pre>	
+220° ×223× Cyclin G1 ribozyme binding site	
k400 × 2634 gatteetgte taaaatgaa	19
<pre>&amp;210 + 2635 &amp;211 + 19 &lt;212 + DNA &lt;213 + Homo sapiens</pre>	
-D200- SD23 - Cyclin G1 ribozyme binding site	
k400% 2635 tidotgtota aaatgaagg	19
+C10+ 1636 +C11+ 19 +C11+ DNA +C11+ Homo sapiens	
81200- 81238 Cyclin Gl ribozyme binding site	
-400% 1636 aatgaaqqta cagoocaag	19
+0.100+ 1.637 +0.11+ 14 +0.12+ DNA +0.13+ Homo sapiens	
::220°-	

-1.223 Dyclin 31 ribozyme binding site	
-:400 + 1637 baadbabbtt gggtgtgtt	19
<pre>HU10+.638 HU11+19 HU12+ DNA HU13+ Homo sapiens</pre>	
+:220 + +:223 + Cyplin G1 ribozyme binding site	
-:400 + 1638 tgggtgtgtt ggactgago	19
<pre>%210 + 0639 %211</pre>	
<pre>&lt;2008 &lt;2008 &lt;2008 </pre>	
<pre>&lt;400 + 2639 tgagotgott ttatttggc</pre>	19
+0210 × 20440 +0211 × 19 +0212 × 9MA +0213 × Hemo sapiens	
+:27.0 + +:27.0 + +: Syclin G1 ribozyme binding site	
H40% 2040 Gagotgottt tätttggot	19
+0010 + 0641 +0010 + 10 +0010 + DNA +0010 + Homo sapiens	
+02160+ +02070+ Cyclin G1 ribozyme binding site	
<pre>+:400:- 1+41 aget@et*tt atttggetg</pre>	19
+210:- 2642 +2115-19	

H212 + ENA H213 + Homo sapiens	
<pre>2220. 2223. Cyclin G1 ribozyme binding site</pre>	
-1400 + 1642 gotgotttta titggotgt	19
+0010 + 0043 +0011 + 19 +0010 * CNA +0013 * Home sapiens	
<pre><pre><pre><i2230 <="" pre=""> <pre></pre> <pre></pre> <pre>Cyclin G1 ribozyme binding site</pre></i2230></pre></pre></pre>	
<400 × 2643 tgcttttatt tggctgtaa	19
<pre>:C10 + 2644  :C11 &gt; 19  <c10> DNA :C13 &gt; Homo sapiens</c10></pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyclin G1 ribozyme binding site</pre>	
<pre>&lt;400% 1644 gettttattt ggetgtaaa</pre>	19
<pre>#210% 2645 &lt;211% 10 &lt;0110% PNA #2013% Homo sapiens</pre>	
+:1230- -:1230- Cyclin G1 ribozyme binding site	
kuigactyta aaatcaata	19
+0.100 2x46 +0.110 19 +0.110 DMA +0.1130 Home sapiens	
+:210:- +:213:- Cyclin G1 ribozyme binding site	
√400: 2646	

otytaaaato aatagaaga	19
<pre>#2100 2647 #211 + 19 #212 + DNA #213 + Homo sapiens</pre>	
+0000 + +0003 + Cyplin G1 ribozyme binding site	
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-:4005-2649 atqtoccatt ggcaactga	19
+02108 D650 +02108 19 +02108 DMA +02138 Homo sapiens	
<pre>+:D200 +:D230- Cyclin G1 ribozyme binding site</pre>	
+0400% 1650 caactgantt gatoogaat	19
+02100 0x51 +0310 19 +0310 DNA	
<213> Homo sapiens	

<pre>%230 \ %233 \ Typlin G1 ribozyme binding site</pre>	
(400 + 2651 tgacttgatc cgaataagt	19
#210 + 2052 #211 + 14 #212 + 5NA #213 * Homo sapiens	
<pre>&lt;2.00 + &lt;2.03 + Gyolin G1 ribozyme binding site</pre>	
k400> 2652 gatoogaata agtoaatat	19
<pre>&lt;210&gt; 2653 &lt;211&gt; 19 &lt;210&gt; DNA &lt;213&gt; Homo sapiens</pre>	
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-:400 > 1€54 taaqtosata taggittac	19
+0216 > 0655 +0211 + 19 +0212 > TNA +0213 > Homo sapiens	
-002005 -01230 Cyclin G1 ribozyme binding site	
राई0ाल- १६५५ agtowatata ggtttacgg	19
321fr 2656	

-C.11 · 19	
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REAR - Cyclin G1 ribozyme binding	site
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aatataggtt tacggtttc	1 3
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<pre>&lt;.123&gt; Cyclin G1 ribozyme binding</pre>	site
<400 > 2657	
atataggttt acggtttca	19
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K210 ≥ 2658	
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<(400.+ 2658	
tataggtita eggttteag	19
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CLIC DNA	
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+:400:+ 1:659	
gittacqqtt tcagacttg	19
H210H 1660	
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-2115- Homo sapiens	
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<pre></pre> <pre>&lt;</pre>	site
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(400 + 266) httw:gytth cagacttga	19
0210 + 2661 0211 + 19 0212 + DNA 0215 + Home sapiens	
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<pre>04000 1661 ttaeggtttd agacttgat</pre>	19
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0400 + 2662 titoagaott gatgagaat	19
+0.10 × 0.663 +0.11 + 19 +0.213 + DMA +0.213 + Homo sapiens	
<pre><pre><pre>&lt;2200</pre> <pre>&lt;2230 Cyclin G1 ribozyme binding site</pre></pre></pre>	
-(400)- 1663 qqaaaaqatt gtattggag	19
+00100 0.004 +00110 10 +00110 PNA +00130 Homo sapiens	
-miles -miles Cyclin Gl ribozyme binding site	
-:4001664 aaadatidfa ttggagaag	19
+0.100+.769 +0.110+.19 +0.110+.19 +0.110+.19 +0.110+.19 +0.110+.19	

<pre>&lt;:2.23 * Cyclin G1 ribozyme binding site</pre>	
-:400 + 2665	
agattgtatt ggagaaggt	19
+210 + 2666	
KIII + 1M	
<pre><d12 +="" dna<="" pre=""></d12></pre>	
-1213 - Homo sapiens	
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and y system of the segment between greating of the	
+:400 × .7666	
aaqqtgtqtt ggaaagtca	19
+:2.10 × :2667	
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<pre>&lt;213&gt; Homo sapiens</pre>	
<220 \( \text{\text{220}} \)	
+223% Cyclin G1 ribozyme binding site	
- Q10 - Q457	
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ttggaaagtd aaagetact	19
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ttggaaagtd aaagdtact -:210 2668	19
ttggaaagto aaagotact +210+2668 +211+13	19
ttggaaagto aaagotact -:210 2668 -:211 19 -:212> DNA -:213> Homo sapiens	19
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ttggaaagto aaagetact -210 - 2668 -211 - 13	19
ttggaaagto aaagetact -:210 - 2668 -:211 - 19 -:212> DNA -:213> Homo sapiens -:223> Cyclin Gl ribozyme binding site	
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tiggaaagto aaagotact -2102668 -21119 -212>-DNA -2013>-Homo sapiens -2023Cyclin G1 ribozyme binding site -4400-2668	
ttggaaagtd aaagctact  -0100668 -0110-19 -0105 DNA -0105 Homo sapiens  -0106-2668 -0106-2668 -0106-2669 -0110-19 -01105 DNA -01105	

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-11 13:00	Homo sapiens		
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-122(1)	Q.,_3 /_ Q1	3-2-31	
. 223-	Cyclin G1 ribczyme	prinaring	site

-:400 · 2674	
optitionality ofgonacty	19
H210 + 2675	
H211 + 19	
+0.1.0 + DNA	
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+:220 +	
<223 - Cyclin G1 ribozyme binding site	
+(400 + 2675	
tttbaattte tgcaactgt	19
+(210 + 2676	
K211 + 19	
K2127 DNA	
<213 Momo sapiens	
K220 ×	
<pre>&lt;223&gt; Cyclin G1 ribozyme binding site</pre>	
+:400 + 2676	
tgcaactgta ttattcact	19
+0.10 + 2677	
+1211 + 19	
HELE FINA	
Hill3: Hemo sapiens	
<(200)>	
K223S Cyclin G1 ribozyme binding site	
+14001+ 2x77	
daadtgtatt attoactop	19
H110N 0678	
+:211:- 19	
+:0100 DNA	
-: 213: Homo sapiens	
-00280 Cyclin G1 ribozyme binding site	
- 100 - 1 - 70	
-:4000- 2678	10
aactqtatta tteacteet	19
-C 100 . 679	
-0.100 (n/g) -0.110 (g)	
H212> DNA	
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-:213 · Homo sapiens	
x229 ·	
<223 - Syclin G1 ribozyme binding site	
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orgitatists captocite	19
+:010 + 0680	
<211 / 19	
<210 - DNA	
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Cliss fond sapiens	
killius	
-MMBA Cyclin Gl ribozyme binding site	
k(406/s) [1682]	
tfdactrott caagagaac	19
K210% 0683	
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edante	
4007 Cyclin G1 ribozyme binding site	
thactocttc aagagaact	19

+:210 + 0.684 +:211 + 13 +:212 + DNA +:213 + Homo sapiens	
+0.00 + +0.00 - 4.00 -	
-:400 - 2084 Aagagaactt godacttga	19
+010 + 0005 +0110 + 19 +0210 + DNA +0213 + Homo sapiens	
H2208 H223> Cyclin G1 ribozyme binding site	
-:400 - 2685 cttqccactt gaaaggaga	19
H210 × 2686 H211 × 19 H212 × DNA	
+1213 + Homo sapiens +1210 +	
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4400% 1686 aggagaaata goattaatt	19
+110 + 1687 +111 + 19 +111 + DNA	
+2213 + Homo sapiens	
+:220.+ +:223:- Cyclin Gl ribczyme binding site	
-:400:- 1687 aaatagcatt aattttgaa	19
+21(0+ 2488 +221(+ 19 +2213+ DNA +2213+ Homo sapiens	
·:220:	

<pre>&lt;0.23 • Cyclin 31 ribozyme binding site</pre>	
<pre>0400 + 0688 agrantmatt ttgaaagac</pre>	19
<pre>%210 + 2689 %211 + 19 %212 + DMA %213 * Homo sapiens</pre>	
·:()(1)() >	
-0.03 > Cyclin G1 ribozyme binding site	
H400 × 2689 gcattaattt tgaaagact	19
+010> 0690 +011> 19 +010> ENA +013> Homo sapiens	
-220 > -223 > Cyclin Gl ribozyme binding site	
0400% 2690 mattaattiti gaaagadta	19
0210 × 2691 0211 × 19 0212 × DNA 0213 × Homo sapiens	
<pre>&lt;2208 &lt;23 * Cyclin G1 ribozyme binding site</pre>	
-0400+ 0691 tgaaagasta gaagotoaa	19
HILLON 1692 HILLON 19 HILLON ENA HILLON Homo sapiens	
01700 01750 Cyclin Gl ribozyme binding site	
(400) lr92 ofadaagoto aactgaagg	1.9
<210:- 2693 <211:- 19	

H212 + DNA H213 + Hamp sapiens	
+(22) + (223 + Gyolin G1 ribozyme binding site	
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+1010 + 0694 +1011 + 19 +0110 + DNA	
-1113 - Homo sapiens	
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k400 - 2694 qoatqtoatt goaggatoa	19
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<210% 2696 <0011% 19 <0012% 0MA <0015% Homo sapiens	
*C20*	
+00030 Cyclin G1 ribozyme binding site	
त्र्विण्ट प्रतिकृति । daggateata ttttetaaa	19
-0.10% 1697 -0.21% 19	
+C2120+ DMA +C2130+ Homo sapiens	
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<4400> 2697	

ggatcatatt ttotaaagc	19
<pre>0010 + 0698 0011 + 19 00108 DNA 0013 + Homo sapiens</pre>	
HD20 > HD23 + Myclin G1 ribozyme binding site	
-:400 - 1698 Matcatattt totaaagoa	19
+010> 0699 +011> 19 +010> DNA +013> Homo sapiens	
<pre><dustrian <a="" href="https://www.commons.com">ccccccccccccccccccccccccccccccccccc</dustrian></pre>	
-400> 1699 atcatatttt ctaaagcaa	19
<pre>%210 &gt; 2700 %211 &gt; 19 %212 &gt; EMA %213 &gt; Homo sapiens</pre>	
<pre>&lt;2200* &lt;223* Cyclin G1 ribozyme binding site</pre>	
+:4005 1700 toatatitto taaagcaaa	19
+210> 2701 +2115 19 +2125 ENA +2135 Homo sapiens	
+0.000+ +0.003+ Cyclin G1 ribozyme binding site	
0400% 2701 atatitiota aagcaaagc	19
+02100+0702 +0210+04 +02120+DNA +02130+Homo sapiens	

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<pre>2: Cyclin 31 ribozyme binding site</pre>	
· 400 · 2702	
graaageett otgtgttgg	19
+210 + 2703	
+211 + 19	
×1113 × DNA	
+213 - Homo sapiens	
rilao -	
<pre>&lt;.123 - Cyclin G1 ribozyme binding site</pre>	
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caaaqootto tgtgttggo	19
3224,00 09290099	
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K215 Hemo Sapiens	
k 220%	
-223 · Cyclin G1 ribozyme binding site	
- 400 × 2705	
tattageatt gtetateat	19
· 210 · 2006	
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<pre>&lt;213 Homo sapiens</pre>	
+ 023 + Cyclin G1 ribozyme binding site	
· 400 · 2706	
tygeattgte tateattge	19
·.210 · 2707	

+1211 + 13 +1212 + DMA +1213 + Homo sapiens	
<pre>&lt;:220 - &lt;:223 - Cyclin G1 ribozyme binding site</pre>	
<pre>&lt;4(0) + 3707 grantgtota toattgcat</pre>	19
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#210% 2709 #211> 19 #212> DNA #213> Homo sapiens	
+MMLO> +MMLO> Cyclin G1 ribczyme binding site	
+400> .709 gtotatratt goattagag	19
+210> 2710 +211> 19 +211> DNA +211> Homo sapiens	
+220> +223> Cyclin Gl ribezyme binding site	
+:400> 2010 toattgoatt agagatoca	19
+210> 7011 +211> 13 +212> ENA +213> Homo sapiens	
+:220:-	
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0400 + 0711 Cattgoutta gagatocaa	19
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+400> 2712 attagagato caagcacag	19
<210> 2713 <211> 19 <212> DNA <213> Homo sapiens	
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+400 + 1715 tqtawaqtta acagaagga	19
+ 210 + 2316 + 211 + 19 + 212 + EMA	

-CRAD	
्र400 + 2716 चत्रवक्ष्मgaata gaatgtott	19
-0.19 + 2717 -0.211 + 19	
RC113 - DMA RC13 - Homo sapiens	
+220.4 +223.4 Cyclin G1 ribozyme binding site	
- (4005 1717	19
atagaatgto ttoagaaab	19
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TOTAL TOTAL	
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aqaatgtott cagaaacat	19
010 010	
+2108 2719 +2118 19	
KOLEN DNA	
RC138 Homo sapiens	
H020N	
-0220 Cyclin Gl ribozyme binding site	
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-00130 Homo sapiens	
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cagaaacatt ccaagataa	19

H210 + 0721 H211 + 19 H212 + DNA H213 + Homo sapiens	
+220+ +223+ Cyclin G1 ribozyme binding site	
H400 - 2721 agaaacatto caagataaa	19
<pre>-HH10 + 17HH -H11 + 13 -H110 + DMA -H110 + Home sapiens</pre>	
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+0210> 0703 +0211> 19 +0210> DNA +0213> Homo sapiens	
+2200 +223 + Cyclin 31 ribozyme binding site	
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+1710 + 1714 +1211 + 19 +1211 + DNA +1213 + Homo sapiens	
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+210+1725 +211-14 +212-DNA +213-Homo sapiens	
HAMICH HAZZBF Cyclin Gl ribozyme binding site	

+:400 + 0725 tatgacette tggcaagag	19
+1210 + 2726 <211 + 19 <213 + DNA <213 + Homo sapiens	
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agottg:ato casatgttt	19
+2108 2704 +2119 19 +2110 DNA +2110 Home sapiens	
+.220S	
+223> Cyclin G1 ribozyme binding site	
<pre><i4000 *ccaaatqtt="" .72.*="" pre="" taactgaat<=""></i4000></pre>	19
+:2108 :2730 +:211:-13 +:212> DNA	

<pre>&lt;:213 · Homo sapiens</pre>	
<pre>-::20 + -::223 - Cyclin G1 ribozyme binding site</pre>	
+:400 + 2730 obaaatgitt aabtgaata	19
H210 H 2731	
H211 + 19	
HU10 + DNA HU13 + Humb sapiens	
+:229×	
<pre>&lt;223&gt; Cyclin G1 ribozyme binding site</pre>	
-:400% 2731	
caaatgttta actgaatat	19
<pre><i310 *="" 2732<="" pre=""></i310></pre>	
+1211 × 19	
<pre>S2128 DNA &lt;1.138 Homo sapiens</pre>	
* nomo sapiens	
<10.20 ×	
<pre><pre><pre><pre><pre></pre></pre></pre><pre><pre><pre></pre></pre></pre><pre><pre></pre></pre><pre></pre></pre><pre></pre></pre> <pre></pre> <pre><td></td></pre>	
\$400 + 2732	
taactgaata tteatcaaa	19
ki210 × 2733	
S2115 19	
PODICE DNA	
+2213 · Homo sapiens	
-11120 s	
+00.03 + Cyclin G1 ribbzyme binding site	
×4400 + 2733	
actgaatatt catcaaata	19
+1210 - 2734	
+12118 19	
+DID-DMA	
Home sapiens	
+111,00+	
+22.30 Cyclin G1 ribozyme binding site	
<400> 2734	
ctgaatatto atcaaataa	19

+1210 + 2735 +1211 + 19 +1312 + DNA +1213 * Homo sapiens	
+:220 + +:223 * Cyclin G1 ribozyme binding site	
- 400 - 2735 aatattoato aaataagtg	19
+210 + 3736 +211 * 19 +212 * DNA +213 * Homo sapiens	
<pre>&lt;220* &lt;0.03* Cyclin G1 ribozyme binding site</pre>	
<pre>&lt;4008 2736 toatcaaata agtgttoca</pre>	19
+1210 × 2737 +2211 × 19 +2213 × Homo sapiens	
<pre>&lt;2210&gt; &lt;2135 Cyclin G1 ribozyme binding site</pre>	
-:400% 2737 aataagtqtt ccaaaccaa	19
+Thics 2738 +Thics 19 +Thics DNA +Thics Homo sapiens	
+2234> +2234+ Cyclin G1 ribozyme binding site	
+400: 2738 ataagtgtto caaaccaaa	19
+210+2739 +211+1+ +211.+ DMA +213+ Homo sapiens	
+:2200	

$\pm (223 + 3 \text{yolin 31 ribozyme binding site})$ $\pm (400 + 1739)$	
abbasat att cagaagttg	19
H210 + 1740 H211 + 19 H312 + DNA H313 + Hamb sapiens	
CDDGS Clock Cyclin G1 ribozyme binding site	
k400+ 1740 ccaaatditc agaagttga	19
<210> 2741 <211> 19 <212> DNA <213> Homo sapiens	
<2200> <2210> Cyclin G1 ribozyme binding site	
<pre>&lt;4000-1741 ttdagaagtt gaaatggat</pre>	19
<pre>32100 2742 42110 19 42128 DNA 82138 Homo sapiens</pre>	
k00000 k00000 Cyclin G1 ribozyme binding site	
-:4005-2742 gaaatggatt gtttotggg	19
SUM108 .743 -UN110 14 -UN120 DNA -UN130 Home sapiens	
-mines -mines Cyclin G1 ribozyme binding site	
+:400:2743 atggattgtt totgggogt	19
+12101- 2744 +12111- 19	

·C12 · DNA ·C13 · Homo sapiens	
+23C+	
- 203 - Cyclin G1 ribozyme binding site	
+490× 2744	
tygattyttt ctgggcgta	19
- 210 > 2745	1.9
- 311 + 19	
· 112 · DNA	
:013/ Homo sapiens	
+ 1202	
<pre>&lt;223&gt; Cyclin G1 ribozyme binding site</pre>	
<400> 2745	
agattgttto tgggogtao	19
<210> 2746	10
<211> 19	
<pre>&lt;212&gt; DNA</pre>	
<pre>&lt;213&gt; Homo sapiens</pre>	
<200>	
<pre>&lt;223&gt; Cyclin G1 ribozyme binding site</pre>	
<400> 2746	
totgggegta otgcaogge	1.0
<210 > 2747	19
<211 > 19	
<pre>&lt;110&gt; DNA</pre>	
C13> Homo sapiens	
H220%	
SDDF - Cyclin G1 ribozyme binding site	
H400× 2747	
Cacqgcaatt gaagcatag	19
+110 × 2748	13
+311× 19	
1012 > DNA	
· 113 · Homo sapiens	
+216+	
- 2003 - Cyclin Gl ribozyme binding site	
< 400 × 2748	

ttgaagcata gctactaca	19
+010 × 0749 +0.11 + 19 +0.11 > DNA +0.13 > Homo sapiens	
-020. -023- Cyclin G1 ribozyme binding site	
+400 + 2749 agbatagota otacagaat	19
+:010 + 0750 +:011 + 10 +:010 + DNA +:013 + Homo sapiens	
<220× <223> Cyclin G1 ribozyme binding site	
0400% 1750 atagotanta dagaataan	19
+210 + 2751 +211 + 19 +212 + DNA +315 + Homo sapiens	
-1.10 - -1.13 - Cyclin G1 ribozyme binding site	
8400: 2751 otabagaata actbacctt	19
HC190- 1752 HC110- 19 HC110- DNA HC130- Homo sapiens	
କ୍ଷିଥିତ Cyclin G1 ribozyme binding site	
04005-0752 agaataacto accttocaa	19
01.100-1753 01.110-13- 02.120-DNA 02.130-Homo samiens	

÷:22n →	
<pre>&lt;:223 - Cyclin G1 ribozyme binding site</pre>	
(4)0 + 2753	19
aactdabott ccaacaatt	1 2
KC10 + 2754	
H211 + 19	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
-1113- Homo sapiens	
+(220 +	
不記2子・Cyclin G1 ribozyme binding site	
8(4(0) × 2.154	
actinacette caacaatte	19
4	1.
K210> 2755	
<211 > 19	
HO10> DNA	
<213 → Homo sapiens	
K220>	
<pre><d23> Cyclin G1 ribozyme binding site</d23></pre>	
<400→ 2755	
tocaacaatt ootgaaatg	19
Issaacaace Cocyadacy	15
H210> 2756	
-C211+-19	
H212 > DNA	
-213 - Romo sapiens	
<220 \	
<pre></pre> <pre></pre> <pre></pre> <pre>Clid - Cyclin G1 ribozyme binding site</pre>	
(400 + 1756)	
daadaatto otgaaatgg	19
soud add teo etgada tyg	1,
d2100+ 1757	
<2110-19	
COLOR DNA	
+22130- Homo sapiens	
KILOUCH	
00030 Cyclin G1 ribozyme binding site	
	19
tgaaatqgto oottaactg	13
210, 275	

·:210:- 2758

·:211 · · ·:.:12 ·		
-1213·	Homo sapiens	
-1220-		
-12.23 →	Cyclin G1 ribozyme binding site	
-(400 ·		
atgqt(	postt aastggatt	1 9
4210 -		
+1211 ×		
·:212 · -:213 ·	Homo sapiens	
<:220 ×		
	Cyclin G1 ribozyme binding site	
<400>	·75a	
	ectta actggatta	1 9
-		
<2100 <211>		
·2128		
	Homo sapiens	
	The state of the s	
<:220 s		
<2223%	Cyclin G1 ribozyme binding site	
<400 >		
taacto	ggatt attacagca	19
4216×	2761	
C211>		
·1712 ·		
·10135	Homo sapiens	
-1220-		
HI23 +	Cyclin G1 ribozyme binding site	
-:400-	2761	
aactigg	gatta ttacagcac	19
·1210.·	2762	
.:211:-		
12:-	IMA	
-01235	Homo sapiens	
·052001+		
	Cyclin G1 ribozyme binding site	

.

H400 - 0760 Htt#att acagcacca	19
<pre>&lt;210 + 0763 &lt;2211 + 19 &lt;2310 + DNA</pre>	
Homo sapiens	
+:210 + +:	
R400+1763 tqqattatta dagdaddaa	19
H010 - 0764 H011 - 19 H010 - DNA	
<pre>&lt;7.13&gt; Homo sapiens</pre>	
<pre>&lt;223 &gt; Cyclin G1 ribozyme binding site</pre>	
<pre>P400 = 1764 Caaaaaactt ototgaago </pre>	19
K210 + 2765 H211 + 19 H212 + ENA	
+213 + Homo sapiens	
<pre>-:220 - &lt;223 - Cyclin G1 ribozyme binding site</pre>	
4400 × 2765 aaaaaaatto totgaagoo	19
+1210 + 0766 +1211 + 19 +1212 + DMA	
+1713 Homo sapiens	
-TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- - -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -TTDO- -	
+(400)+ 2766 asaaottoto tgaagoott	19
<pre></pre>	
<pre> <pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> </pre> <pre> <pre> </pre> <pre> </pre> <pre> <pre> </pre> <pre> </pre> <pre> </pre> <pre> </pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre> <pre> <pre> <pre> </pre> <pre> <pre> </pre> <pre> <pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	

<pre>&lt;220 -  &lt;223 - Oyolin G1 ribozyme binding site</pre>	
0400 + 2767 obgaagoott totocadaa	19
H210 + 2768 H211 + 14 H212 + DNA H213 + Homo sapiens	
+MMLO+ +MMLO+ Cyclin G1 ribozyme binding site	
k400 + 2768 tgaagoottt otooacaac	19
<pre>%310 / 3769 %311 / 19 %312 + DNA %313 / Homo sapiens</pre>	
<pre>8.220 + 8.223 + Cyclin G1 ribozyme binding site</pre>	
<pre>0400 2769 gaageoffte topacaace</pre>	19
<pre>&lt;210 + 2770 &lt;211 + 19 &lt;212 + DNA &lt;213 + Homo sapiens</pre>	
<pre>s2200 - c2223    Cyclin G1 ribozyme binding site</pre>	
k400 1770 agostitoto cacaaccii	19
+0210+ 0771 +0210+ 14 +0210+ DNA +0213+ Homo sapiens	
HURON Cyclin G1 ribozyme binding site	
-:400:- 2771	19

```
<210 → 2772
-1211 + 19
-1.111 · DNA
+1213 + Homb sapiens
HAMAB: Dyclin G1 ribozyme binding site
<14au + 2772</p>
                                                                      19
baabhttatt ctatggatt
3210 + 2773
\pm 1211 \pm 13
-1210 - ENA
+:213 > Hemo sapiens
4:220a
<!223> Cyclin G1 ribozyme binding site
<400> 2773
aaddttgtto tatggattd
                                                                      19
K2109-2774
H2118 19
4212 - DNA
+213> Homo sapiens
-1220%
\pm 2.33 \pm \text{Cyclin} Gl ribozyme binding site
H400% 2774
                                                                      19
cottattota tggattoca
→:210÷ 2275
\pm 1211 + 19
H2125 DNA
-00135 Homo sapiens
</p
-(4000 - 2275)
totatgqatt ccataatgt
                                                                      19
-01100-1776
000110-19
HOLLONA DNA
Homo sapiens
(1.2.2.1)
<:223> Cyclin G1 ribozyme binding site
```

.

(40) - 2776 Statagatts cataatgtt	19
+1210 + 1777 +1211 + 19 +1211 + DNA +1213 + Homo sapiens	
+220 + +223 + Cyclin G1 ribozyme binding site	
+400 + 2777 qqattobata atgttacaa	19
<pre>#210 + 2778 +211 + 18 +212 + DNA +213 + Homo sapiens</pre>	
+000 + +0003 Cyclin G1 ribozyme binding site	
-:400:- 2778 -ccataatdti acaatggat	19
+010 + 0779 +0115 19 +0105 5NA +0132 Homo sapiens	
+2208 <2238 Cyclin G1 ribozyme binding site	
H4005 2779 cataatgita caatggatt	19
<pre>k2105 2080 k2115 19 k22125 DNA k22125 Homo sapiens</pre>	
SCROS -NRAS Cyclin G1 ribezyme binding site	
4400:- 2780 Adaatgdatt taaagdtat	19
+0100+0081 +0110+19 +0212> DNA	

<pre>&lt;213 · Homo sapiens</pre>	
-1220 -	
4223 · Cyclin G1 ribozyme binding site	
-:400 + :2781	
caatigattt aagetatga	19
<210 + 2182	
K2115-19	
KILII≯ DNA	
<pre><il13> Homo sapiens</il13></pre>	
<0.00%	
<pre>SAMB&gt; Gyolin G1 ribozyme binding site</pre>	
<400> 780	1 0
aatigatita agstatgaa	19
<210> 2783	
<2215-19	
<212> DNA	
<pre>&lt;213 &gt; Homo sapiens</pre>	
<220>	
<223> Cyclin G1 ribozyme binding site	
<400 > 1783	
atttaaqota tgaagotca	19
<2100× 2784	
<211> 19	
K2125 DNA	
<pre>&lt;2130 Homo sapiens</pre>	
V2.26%	
CLASS Cyclin G1 ribozyme binding site	
tatgaagnto aaacatcac	19
01100-1786 	
CLIIN 19	
COLOS DNA	
02130 Homo sapiens	
ent on	
(203) Cyclin G1 ribozyme binding site	
(400): 2785	
stcaaacatc acgagataa	19

<pre><di10 *="" 2786<="" pre=""> <pre><di11 *="" 19<="" pre=""></di11></pre></di10></pre>	
HILL DNA	
<pre><ill13> Homo sapiens</ill13></pre>	
-1.29 -	
<pre>%2003  Oyolin G1 ribozyme binding site</pre>	
-4008-2786	
toangagata agcatgatg	19
HD10+ 1787	
C2115-19	
HILLS THA	
<pre>&lt;013&gt; Homo sapiens</pre>	
<.220>	
<pre>&lt;023&gt; Cyclin G1 ribozyme binding site</pre>	
<400> 2787	
catgatggto toagettgg	19
<210> 1788	
<2115 19	
<210 > DNA	
(213) Homo sapiens	
4220S	
<pre><pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre>&lt;</pre></pre>	
<400 × 2788	
cacaacagta gtcagaagc	19
KD10 S D789	
4011 × 19	
HILL HOUA	
KIIIN Homo sapiens	
HLB# +	
SU230 Cyclin H ribozyme binding site	
040008-2789	
áacaqtaqto agaagoggo	19
001004-2790	
0.110-19	
CLIME DNA	
CM130 Hemo sapiens	
regions.	

-: 223 · Cyclin H ribozyme binding site	
+(400 + 1790	
astyyacett etecagoga	19
ab.jyabbac occoayoya	1./
k(210 + L**)41	
+1211 + 13	
KOID - DNA	
RANTA Homo sapiens	
10 Homes Sapiens	
S.3.20 ·	
HIDDB - Typlin H ribozyme binding site	
.555 - Lystin h libozyme binding site	
H400 + 2791	
ggabottoto cagogagga	19
ggassettsee cagegagga	13
K210 × 2732	
KIII 19	
HILLS ONA	
<pre>&lt;313  Homo sapiens</pre>	
Autor nomo sapiens	
x220 ·	
<pre></pre> <pre>&lt;</pre>	
1223 Gydiin h Tibozyme binding Sice	
+:400× 2792	
	19
accgcaaatt cagatgcaa	17
H210 × 1793	
H211 + 19	
H212+ DNA	
+213 + Homo sapiens	
215 · Monte Sapiens	
K220+	
+22.33 Cyclin H ribozyme binding site	
+(400 × 2793	
boqbaaatto agatgcaaa	19
Jung Suda 1.15 aga egeada	10
H210H 0794	
+2211 + 19	
KO 12 - DNA	
+2150 Homo sapiens	
and the supposition of the suppo	
+111.008	
ANNUAL Cyclin H ribozyme binding site	
+(4)00(x)11194	
ogggaaggtt ottoogaat	19
· , , ,	
-(210)> 2°95	
(211) 19	

tgatscagte tttettgag	19
<210 - 2300	
S211 × 19	
KD12 ← DNA	
<pre>&lt;213 · Homo sapiens</pre>	
-:226 +	
S223 - Cyclin H ribozyme binding site	
227 Sycill in Tibolyme binding Site	
-:400 + 280°)	
atomagnett tettgaged	19
H210 + 2801	
K211 + 19	
K312 + DMA	
<213 - Home sapiens	
<220 ·	
<223 - Cyclin H ribczyme binding site	
<400 + 2801	
todagtottt ottgagodt	19
<210 × 2802	
K211 + 19	
H212 - DNA	
<pre>%213 - Homo sapiens</pre>	
K220 +	
<pre>&amp;2.23 - Cyclin H ribozyme binding site</pre>	
+:400 × 0800	
ocalfottic tigagecto	19
<210% 2803	
H211 + 13	
*121.7 - LMA	
+1113 + Homo sapiens	
H120H	
-:::23: Cyclin H ribozyme binding site	
+:400:+ 0803	
agtotttot: gagootoat	19
+:110+ 1804	
+0110 19	
+C12+ DNA	
H213> Homo sapiens	

<pre><d200 +="" <d203="" gyclin="" h="" pre="" ribozyme<=""></d200></pre>	binding	site	
<pre>+:400 + 3504 ottgagooto atgaagaaa</pre>		1	L 9
+:210 + 2#05 +:311 + 19			
SCHICA DNA SCHICA DNA SCHICA Homo sapiens			
R2209 R223> Cyclin H ribozyme	binding	site	
-:400> 2805	zinaing		
aatgacacto tgcaaatac		1	L 9
K210> 2806 K211> 19			
<pre>K212&gt; DNA K213&gt; Homo sapiens</pre>			
<220>			
-0003> Cyclin H ribozyme	binding	site	
-:400 > 3806		1	<u> </u>
totgoasata otatgagaa		1	. 3
<pre>&lt;210 &gt; 2807</pre>			
H211> 19 H212> DNA			
<213> Homo sapiens			
HAA0>			
+223> Cyclin H ribozyme	binding	site	
-G4005-2807			
quaaatabta tgagaaaag		1	9
-:::10> 1508			
-00115-19			
HOTOS MARK TOTAL			
-0113> Homo sapiens			
- ( <u></u>			
<pre>&lt;:223&gt; Cyclin H ribozyme</pre>	binding	site	
<:400: 1308			
agaaaaaqtt attggaatt	•	1	9
::210: ∠809			

+:211 + 19 +:212 + DNA +:213 + Homo sapiens	
-0220 - -0223 - Cyclin H ribozyme binding site	
ki400 - 0809 gaaaaggitta ttggaatto ——————————————————————————————————	19
<210 - 2810 <211 - 19 <212 - DNA <213 - Homo sapiens	
<220% <223 · Cyclin H ribozyme binding site	
<400 - 2810 assorithatt ggaattetg 1	19
<210 × 3811 <211 × 19 <213 × INA <213 × Homo sapiens	
KDDD - KDDD - Cyclin H ribosyme binding site	
k400 - 2811 tattqgaatt otgttoggt — — — — — — — — — — — — — — — — — —	19
<210% 2812 <211% 19 <211% DNA <uli><li>Homo sapiens</li></uli>	
<2200 	
o4000-2812 attoqaatto tgttoggtg ——————————————————————————	19
07100-2813 07110-14 07110-DNA 07130-Homo sapiens	
-02200 -02200 Cyclin H ribozyme binding site	

<pre>&lt;400 &gt; 0813 gaattotgtt oggtgttta</pre>	19
<pre>:C10* L814 :C11* 19 :C10* LMA :C13* Homo sapiens</pre>	
+0205 +02235 Gyolin H ribozyme binding site	
<pre>&lt;400&gt; 2814 aattotqtto ggtgtttaa</pre>	19
<pre>&lt;210&gt; 2815 &lt;221&gt; 19 &lt;2212&gt; ENA &lt;2213&gt; Homo sapiens</pre>	
-020> -023> Cyclin H ribozyme binding site	
<pre>&lt;400 &gt; 2815 qttoggtqtt taagccagc</pre>	19
+00108 2016 +00118 19 +00128 DNA +00138 Homo sapiens	
-22008 -2223> Cyclin H ribozyme binding site	
-4005-2816 ttoggtattt aagecagea	19
+U10 > I917 +U11 > I9 +U10 > ENA +U13 > Homo sapiens	
-mmg- -mmg- -mmg- Cyclin H ribozyme binding site	
k400 × 2817 toggtgttta agccagcaa	19
<pre> <pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	

<pre><!--220.* ril230 Cyclin H riboxyme binding site</pre--></pre>	
k4000-3818 tgccaagate tgttgtggg	19
+02100+ 2819 <0110+ 19 +0110> DNA <013> Homo sapiens	
<2230 Cyclin H ribozyme binding site	
<400> 0819 aagatotgtt gtgggtacg	19
<pre>&lt;010&gt; 0820 &lt;011&gt; 19 &lt;012&gt; DNA &lt;013&gt; Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyclin H ribozyme binding site</pre>	
<400> 2820 gttgtgggta oggottgta	19
<210> 2821 <211> 19 <212> DNA <213> Homo sapiens	
<pre>sch20b &lt;2238 Cyclin H ribozyme binding site</pre>	
k400N 2921 gqtacgqctt gtatgtatt	19
\$210% 2822 \$211% 19 \$212% DNA \$213% Homo sapiens	
+(223)+ Cyclin H ribozyme binding site	
+(400):- 2822 acqqcttqta tqtatttca	19

<pre>&lt;210 * 3833 &lt;2211 * 19 &lt;212 * DNA &lt;213 * Homo sapiens</pre>	
+220 + +223 + Syclin H ribozyme binding site	
H400+ 3833 Stigtatgta titicaaacg	19
+C10 + D804 +C11 + 19 +C10 > DNA +C13 > Homo sapiens	
<220> <223> Cyclin H ribozyme binding site	
<400% 2824 tgtatgtatt tcaaacgtt	19
<210 > 2825 <211 > 19 <212 > DNA <213 > Homo sapiens	
<pre>&lt;120 &gt; &lt;1223 &gt; Cyclin H ribozyme binding site</pre>	
<400> 2825 gtatgtattt caaacgttt	19
<pre>&lt;210&gt; 2826 &lt;211&gt; 19 </pre>	

<pre>0400 + 1927 ttoquacytt tttatotta</pre>	19
+1210 + 1828 +1211 > 19 +1212 + DMA +1213 + Homo sapiens	
+Quas	
HAMB & Cyclin H ribozyme binding site	
<pre>&lt;400 / 2828 tuasaogttt ttatottaa</pre>	19
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre><c113> DNA </c113></pre> <pre><c113> Homo sapiens</c113></pre>	
+:220 × <223 > Cyclin H ribozyme binding site	
H400> 0809 baaacgttti tatottaat	19
+2105 1830	10
×2218 19	
< 212 % DNA	
+213> Homo sapiens	
< 220 >	
<pre><!--::3--> Cyclin H ribozyme binding site</pre>	
+(400) + (0530)	
aasogttitt atottaata	19
K2108 2831 K2118 19	
HULL OF THA	
Homo sapiens	
KUL 03	
-CCCCC Cyclin H ribozyme binding site	
<4000 2831	
aacutttta tottaataa	19
(C100- 1832	
KUNIN IO KUNDA DNA	
NO. 1 4 4 5 174 TO 1	

Homo sapiens	
+1220 +	
+223 + Cyclin H ribozyme binding site	
+(4)00 × 2832	
equititate ttaataaet	19
<:210 + 2833	
K211 × 19	
HOIDS DNA	
+:213 · Homo sapiens	
K2202	
<223 Cyclin H ribozyme binding site	
4400× 1833	
tittitatett aataactea	19
<2108 2834	
<211 × 19	
SOLDS DNA	
+213 + Homo sapiens	
<:220>	
<pre><dd><pre></pre> <pre>Class Cyclin H ribozyme binding site</pre></dd></pre>	
+3400 × 2834	
ttttatetta ataactcag	19
K010% 0835	
S0115-19	
×212× DNA	
+213 - Homo sapiens	
-120A -	
FIRE Cyclin H ribozyme binding site	
-:400% . 935	
tatottaata actoagtaa	19
+0100+0836	
-(211)- 19	
COLOR DNA	
+31.00 Homo sapiens	
(21.C)	
<pre><!--draw Cyclin H ribozyme binding site</pre--></pre>	
·:400> 2836	
ttaataactc agtaatgga	19

<210 · 03337	
3211 * 1.*	
HOLD NOMA	
<pre>#213 &gt; Homo sapiens</pre>	
<.220 *	
<pre><pre><pre><pre></pre> <pre>Cyclin H ribozyme binding site</pre></pre></pre></pre>	
+:400 + £537	
taadtoagta atggaatat	19
H210 + 1838	
H211 × 19	
HIBID > DINA	
+213> Homo sapiens	
+:220>	
+223> Cyclin H ribozyme binding site	
and a specific in the second s	
+:400 × 0838	
taatggaata toaccocag	19
Jan ggazez esassosag	
÷0010 × 0839	
+0.011 > 19	
HILL > DIA	
H213 Homo sapiens	
all, had supreme	
-0.210 s	
S223 - Cyclin H ribozyme binding site	
y star in transpino brinding bres	
k(400 × 2839	
atggaatato accocagga	19
+0°10 + .7840	
+C 11 + 19	
SOLIC - DNA	
+2213 + Homo sapiens	
Title Heller Dup Lotto	
+10.2 C.+	
-0.23 Cyclin H ribozyme binding site	
The state of the s	
+(400) + (2840)	
coccagnata ataatgoto	19
manag as a sucure good	± 2
+12101+ 1841	
-2210 19	
HULLUS DIA	
H213: Homo sapiens	
Little Homo Dapitono	
+220°2	

<pre>&lt;223 - Cyclin H ribozyme binding site</pre>	
<pre>(40) = 2941 taggatauta atgotoact</pre>	19
H210 + 2842 H211 + 14 H212 + DNA	
+1113 + Homo sapiens	
+00.00+ +00.03+ Cyclin H ribozyme binding site	
kupon k. 1842 aataatgoto acttgtgca	19
<pre>#CD10 &gt; 1843 #CD11 &gt; 19 #CD12 &gt; DNA #CD13 &gt; Homo sapiens</pre>	
<pre>&lt;1.00&gt; &lt;223&gt; Cyclin H ribozyme binding site</pre>	
-400% 2843 atgeteaett gtgeatttt	19
<pre>%010 % 2844 %011 % 19 %012 % ENA %013 % Home sapiens</pre>	
<pre>%220% &lt;220% Cyclin H ribozyme binding site</pre>	
-400 - 1844 ottgtgoatt tittggootg	19
+0210 + 2345 +0210 + 19 +0210 + DNA +0213	
+:000@s	
+0023 - Cyclin H ribozyme binding site	
::40c:- 1845 ttgtqcartt ttggcctgc	19
(1210) - 2946 (211) - 19	

H212 - DNA -113 - Humo sapiens	
<pre><dino. <pre="">chin</dino.></pre>	
k400 + 2846 tatacatttt tggcotgca	19
KC10 + 2847 HC11 + 19 HC12 + DNA HC13 + Hemo sapiens	
<pre>&lt;2205 &lt;2235 Cyclin H ribozyme binding site</pre>	
k400 - 2847 gtgbattttt ggootgbaa	19
<pre>4210 \ 2848 4211 \&gt; 19 4212 \ DNA 4213 \&gt; Homo sapiens</pre>	
+220 + +223 - Cyclin H ribozyme binding site	
0400 × 2848 Otgoaaagta gatgaatto	19
+0010> 0849 +0010> 19 +0010> DNA +0013> Homo sapiens	
<pre><pre><pre></pre> <pre></pre> <pre></pre> <pre>Color:</pre> <pre>Cyclin H ribozyme binding site</pre></pre></pre>	
-:400:- 1849 taqatgaatt caatgtato	19
+1210:+ 12850 +1211:+ 19 +1211:+ DNA +1213:+ Homo sapiens	
<pre><pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><!--</td--><td></td></pre></pre></pre></pre>	
÷:400> 2850	

agatraatto aatgtatot	19
<pre>4010 + 2951 4011 + 19 4012 + DNA 4013 + Homo sapiens</pre>	
HUDDA - Oyolin H ribozyme binding site	
-:400 - 2851 attowatgta totagtost	19
<pre>8010 + 0852 8011 + 19 8012 + DNA 8013 * Homo sapiens</pre>	
<2210 - <223 - Cyclin H ribozyme binding site	
+400> 2852 toaatgtato tagtootoa	19
<pre>&lt;210.0 2853 &lt;2211. 19 &lt;2210.5 DNA &lt;2213.5 Homo sapiens</pre>	
-:220 - -:233 Cyclin H ribozyme binding site	
<pre>&lt;400% 1853 aatgtafota gtootoagt</pre>	19
+.210% 2854 +1.11% 19 +0.11% IMA +0.13% Home sapiens	
-(226)- -(227)- Cyclin H ribozyme binding site	
-:400:- 2854 gtatotagto otoagtttg	19
+0.100+ 0858 +0.110+ 04 +0.110+ DNA +0.110+ Beams samings	

<.2203	
-:223 - Cyclin H ribozyme binding site	
+(400 + 2955	
notagtouto agtitigity	19
-:210 - 2856	
-:::::: 19	
HARLY DNA HARLY Homo sapiens	
K220* K223* Cyclin H ribozyme binding site	
-:400 / 1856	
gtootoagtt tgttggaaa	19
-:210 × 2857	
K211 × 19	
RUIDA DNA RUIDA Homo sapiens	
The Tento Sapiens	
K220 /	
AMM3> Cyclin H ribozyme binding site	
-:400 × 2857	
tootcaqttt gttggaaac	19
+:210 × 2858	
-1211 > 19	
HOID FINA	
*213> Homo sapiens	
H220 K	
+27.3 + Cyclin H ribozyme binding site	
-1400 + 0 = 5 =	
thagttigtt ggaaacctc	19
H210H 1859	
S01115 19	
+MAINS DNA +MAINS Homo sapiens	
KIND OF	
+MM 3> Cyclin H ribozyme binding site	
+(400)+ 2853	
tqgaaacetc cgggagagt	19
K2100-2860	

<pre>%211 * 19 %212 * DNA %213 * Homo sapiens</pre>	
kaaa . Kaaa . Oyalin H ribozyme binding site	
-:400+ 1860 opygagagto otottggad	19
+210 × 2861 +211 > 19 +212 × DNA +213 > Homo sapiens	
<220> <223> Cyclin H ribozyme binding site	
<400> 2861 gagagtooto ttggacagg	19
<pre>:210&gt; 2862 :211&gt; 19 :212&gt; DNA :213&gt; Homo sapiens</pre>	
-2008 -2013: Cyclin H riboxyme binding site	
-4005 1860 gagtoctott ggacaggag	19
+2108 2863 +2118 19 +2118 ENA +213> Homo sapiens	
k220x k220x Cyclin H ribozyme binding site	
::4(pp: 1863 qaaqqoactt gaacagata	19
+00100 0864 +0010 1+ +0010 DNA +0013 Homo sapiens	
H2200 H2200 Cyclin H ribozyme binding site	

-:400 - 2864 tgaabagata otggaatat	19
+0010 + 0865 +0011 + 19 +0013 + DMA +0013 + Homo sapiens	
<pre><pre><pre><pre></pre> <pre><pre></pre> <pre></pre> <pre><td></td></pre></pre></pre></pre></pre>	
-:400% 1865 Tachggaata tgaactact	19
+1210 > 1866 +1211 > 19 +1212 DNA	
+213 > Homo sapiens +220 - +223 > Cyclin H ribozyme binding site	
1400 × 2866 atatgaacta ottottata	19
H210> 2867 H211+ 19 H213> PMA	
<pre>+C213 &gt; Homo sapiens +C220.*</pre>	
<pre><pre><pre><pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><!--</td--><td></td></pre></pre></pre></pre>	
tigaantuntti ottatadag	19
+02100	
+:::20:- -:::23:- Cyclin H ribozyme binding site	
-:400: 2868 antantinti atacagcaa	19
<pre>#C100</pre>	

+3200 +323 - Cyclin H ribozyme binding site	
-(400 + 2869 Stasttita tacagoaac	19
H010 + 0870 H011 > 19 H010 + DNA H013 + Homo sapiens	
<pre><pre>&lt;220 -</pre> <pre>&lt;223 - Oyclin H riboxyme binding site</pre></pre>	
<pre>&lt;400 &gt; 2870 abttottata dagdaadtt</pre>	19
<pre>&lt;210 * 2871 &lt;211 * 19 &lt;212 * DNA &lt;213 * Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223 - Cyclin H ribozyme binding site</pre>	
-:400 - 2871 abagcaactt aattteeac	19
+210+ 2872 +211+ 19 +212+ DNA +213+ Homo sapiens	
+URBS - Cyclin H ribozyme binding site	
+:4008 2872 dagdaactta atttccacc	19
+00100 2873 +00110 14 +001.5 DNA +00130 Heme sapiens	
+Clics +Clic Cyclin H ribozyme binding site	
<pre>&lt;4000:- 287s</pre> caacttaatt tocacctta	19

<pre>&lt;210 + 3874 +0.11 + 19 +0.12 + DNA +0.13 + Homo sapiens</pre>	
<pre>&lt;:220 - &lt;:223 - Gyolin H ribozyme binding site</pre>	
e400 - 1874 aasttaattt ccaccttat	19
+1110 + 2375 +111 + 19 +112 + DNA +113 > Homo sapiens	
<pre><date="10.000"> <date="10.000"> <date="10< td=""><td></td></date="10<></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></date="10.000"></pre>	
<pre>4400&gt; 0875 acttaatttc caccttatt</pre>	19
+210> 0876 +211> 19 +210> DNA +213> Homo sapiens	
<pre><!--200--> <!--2038 Cyclin H ribozyme binding site</pre--></pre>	
H400> 1876 tttscacett attgtccac	19
+:2105 2877 +:2112 14 +:21.0 LMA +:2130 Homo sapiens	
+2160+ +223% Cyclin H ribozyme binding site	
ki400% 2877 ttonachtta ttgtocaca	19
+00100+ 2878 +00108 14 +00100+ DNA +00100+ Homo sapiens	
-C.No	

<pre>&lt;:400 * .878 ppappttatt gtocacaat</pre>	19
<pre>%210 + 2879 %211 + 19 %212 + DNA %213 + Homo sapiens</pre>	
+0.10 + +0.03 + Cyclin H ribozyme binding site	
-1400 - 2879 pottattqto cacaatoot	19
<pre>&lt;210 - 2880 &lt;2211 - 19 &lt;2212 - DNA</pre>	
K213 - Homo sapiens K220 -	
<pre><d2.13 -="" 2880<="" <400="" binding="" cyclin="" h="" pre="" ribozyme="" site=""></d2.13></pre>	
gtocacaato ottacagac +:210 + 2881 +:211 + 19	19
+U.12 + DNA +U.13 + Homo sapiens	
<pre>kDD0 &gt;     Cyclin H ribozyme binding site</pre>	
<pre>-:400 + J881 cadaatoott adagaddat</pre>	19
RC100 - 1882 RC110 - 10 RC120 - DMA	
<pre>+M130: Homo sapiens summary during the single state of the si</pre>	
+11030 Cyclin H ribozyme binding site +140C0 1882	19
adaa cotta dagaccatt -1110-1883 -1111-13	19
+121212 DNA	

4.313 · Homb sapiens	
<pre>0.30 color H ribozyme binding site</pre>	
<pre><i400 +="" 2883="" adagactatt="" pre="" tgagggett<=""></i400></pre>	19
+0210 + 0884 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
RANDO	
+3400x 2884 bagaboattt gagggotto	19
<pre>#210% 2385 #211% 19 #2213% DNA #2213% Homo sapiens</pre>	
<pre>&lt;:220&gt; :223&gt; Cyclin H ribozyme binding site</pre>	
H400 + 3885 ttqaaggatt actaataga	19
+0010% 0686 +0010% 19 +0010% DNA +0013% Homo sapiens	
+.22005 +.2200- Typlin H ribozyme binding site	
+4005-1886 igagggette eteategae	19
+210+ 2887 +211+ 19 +212> DNA +213> Homo sapiens	
+0.200+ +0.2030+ Cyclin H ribozyme binding site	
+:400:- 1887 gggcttcctc atcgactta	19

÷(210)÷ 2888	
+1211 + 1 #	
+012 + 0MA	
+213 · Homo sapiens	
+(2.2.6.2)	
+2223 + Cyclin H ribozyme binding site	
+:400 + 28 <b>8</b> 8	
ottootoato gaottaaag	19
H210 + 2889	
<211 × 19	
HILLS DMA	
-213> Homo sapiens	
215 Home Sapiens	
H2208	
<pre>&lt;223 * Cyclin H ribozyme binding site</pre>	
+400 × 2889	
toatogaett aaagaceeg	19
<211 • 19	
H212 N DNA	
<pre>+C13 - Home sapiens</pre>	
(0.00 + )	
·M.3 · Cyclin H ribozyme binding site	
K400+- U890	
categactta aagaccege	19
H210 + 2891	
H211 + 19	
KANION DINA	
HD17: Homo sapiens	
(020)	
+023 + Cyclin E ribozyme binding site	
-:400:-1:8 <b>91</b>	
agaccogota toccatatt	19
αφανωσησία τουσατατί	19
42100-1892	
42110 19	
GUIDE DNA	
40130 Homo sapiens	
<2200x	

<pre>&lt;:223 - Cyclin H ribozyme binding site</pre>	
-::400 + 28 <b>9</b> 2	
abbogotato ocatattgg	19
-:210 + 28 ≠3	
H211 × 19	
HO12 + ONA	
<pre><!--!!!3 * Homo sapiens</pre--></pre>	
H220 F	
<223 · Oyolin H ribozyme binding site	
+:400 + 2893	
	19
ctat socata (ttggagaat)	13
<210 · 2894	
<2115-19	
K21D> DNA	
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< 320 •	
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<400 + 2894	
atobbatatt ggagaatoo	19
a. D. D. L. T. C. G.	
+:210 + 2895	
HOLD + 19	
SA1.2 - DNA	
x113 - Homo sapiens	
-K226 K	
+223. Cyclin H ribozyme binding site	
+(400)+ 1895	
tiggagaato cagagattt	19
-1210h . 596	
+:2115-19	
+(212)+ DMA	
+2135 Homo sapiens	
s(110)s	
+: LLS: Cyclin H ribozyme binding site	
+:400> 2896	
tocagagatt ttgaggaaa	19
<:210:- 2897	
<211> 19	

<pre>H212 + DNA H0mo sapiens</pre>	
+:220 + +:222 +: Oyolin H ribozyme binding site	
-:4J0 × La97	
ssadagattt tgaggaaaa	19
+0210 × 0898 +0211 × 19	
HILLS DNA HILLS Homo sapiens	
<pre>child color</pre>	
H:400 - 1898 Baqaqatttt gaggaaaac	19
	10
+:210 + 2899 +:211× 19	
+212: 5NA	
+1135 Homo sapiens	
+220> +2238 Cyclin H ribozyme binding site	
+:400× 0899 ofgatgactt tottaatag	19
	12
<pre>&lt;2105 3900 </pre>	
SCIC DNA	
+D13 + Homo sapiens	
-:11:0 s	
+223 + Cyclin H ribozyme binding site	
+:40C:- 190G	
tqatqantti ottaataga	19
+D103 0901	
+0.110 19 +0.170 DNA	
HILL SHAN	
+11mror-	
+2233 Cyclin H ribozyme binding site	
+:400> 2901	

gatgacttt: ttaatagaa	19
<pre>#210 + 0902 #211 + 19 #212 + DNA #213 * Homo sapiens</pre>	
<pre><dino :<="" td=""><td></td></dino></pre>	
<pre>kqu0 &gt; 1902 tgactttott aatagaatt</pre>	19
<pre>&lt;110 &gt; 2903 &lt;211 : 19 &lt;212 &gt; ENA &lt;213 : Homo sapiens</pre>	
<pre>&lt;220 &gt; &lt;223 &gt; Cyclin H ribozyme binding site</pre>	
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<pre>K210 &gt; 2904 K211 &gt; 19 K212 &gt; DNA K213 &gt; Homo sapiens</pre>	
-220. -223: Cyclin H ribozyme binding site	
+400+ 2904 tttottaata gaattgoat	19
+02100+ 2905 +0210+ 19 +02105 DNA +00130 Homo sapiens	
+0000+ +0023+ Cyclin H ribozyme binding site	
-04000- 2005 taaragaatt goattgaog	19
+00.100+ 0.406 +00.100+ 0.4 +00.100+ 0.NA +02.130+ Homo samiens	

H220 · Haribozyme binding site	
(400 + 2906 quattgoatt gaoggatgo	19
H210 - 2907 H211 - 19	
HODIA - DNA HOMO sapiens	
RIBBON Republic Republic Republic Ribbons Republic Republ	
-(400% 2007 adggatgett adottttat	19
<pre>&lt;210&gt; 2908 &lt;221&gt; 19 &lt;210&gt; DNA &lt;213&gt; Homo sapiens</pre>	
<pre><dmio <="" pre=""> <pre><dmio <="" pre=""> <pre>&lt;2.33 </pre> <pre>Cyclin H ribozyme binding site</pre></dmio></pre></dmio></pre>	
<400> 2908 oggatgotta octtttata	19
<pre>&lt;210 + 0909 &lt;211 &gt; 19 &lt;212 &gt; DNA</pre>	
<213 Homo sapiens	
<pre>0.003 Cyclin H ribozyme kinding site</pre>	
k400 × 0303 igottaccit tiatacaca	19
<pre>&lt;2109 2910 &lt;2210-19 &lt;22128 DMA</pre>	
+2135 Homo sapiens	
+2200+ +2230+ Cyclin H ribozyme binding site	
+:40(:+ z::10) qcttaccttt tatacacac	19
<2210>-2311	

-:::11 · 19	
FILITO - DNA	
+2213 · Homo sapiens	
The transfer of profits	
+320×	
+0023 Cyclin H ribozyme binding site	
+400 + 2911	
ottapottti atacacacc	19
H010 × 0912	
×0.11 + 19	
COLOR DNA	
+1.13 · Home sapiens	
The monte supreme	
<2220 >	
+DDB+ Cyclin H ribozyme binding site	
₹400≥ 0912	
ttacettita tacacacet	19
<210 × 2913	
<211> 19	
4212 × DNA	
<pre>&lt;2130 Homo sapiens</pre>	
ALIST HOME Suprofic	
K2208	
<pre>*CL3 * Cyclin H ribozyme binding site</pre>	
-:400 × 2913	
accttttata cacaccttc	19
<0.10 ± 2914	
2011 - 19	
KO10 + DNA	
-0013 - Homo sapiens	
•	
+1200 ·	
-:223: Cyclin H ribozyme binding site	
.225. Cyc.in in Tibozyme binding Site	
(1AA) 104.1	
+:400:+ ::2914	1.0
tadadadett occaaattg	19
<pre>+0100+ . 915</pre>	
-02115-19	
ROTOR DNA	
+213+ Homo sapiens	
K21.07	
<223> Cyclin H ribozyme binding site	
but of other in thropling chief	

43400+ 2915 abadacutto deaaattge	19
H010 + 0916 H011 + 19 H010 > DNA H013 + Homo sapiens	
+0020 + +0023 + Cyclin H ribozyme binding site	
4400 - 1916 ttoobaaatt goodtgast	19
<pre><c10 -="" 1917="" <c211=""> 19 <c112 -="" <c113="" dna=""> Homo sapiens</c112></c10></pre>	
<pre>&lt;0.20&gt; &lt;0.03 - Cyclin H ribozyme binding site</pre>	
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k:400% 2918 antyddatt: tatdtagtg	19
+00108 (391) +00118 (39 +00128 DNA +00130 Homo sapiens	
+00000 +00000 +00000 Cyclin H ribozyme binding site	
ed@color=2919 otgocatttt atotagtgo	19
+0010 + 1 +20 +0011 + 1 + +0011 + DNA +0013 + Homo sapiens	

AMAGA Cyclin H ribozyme binding site	
-:400 + :3920	
tipoattita totagtgod	19
H110 + U921	
K211 × 19	
H212 + DNA	
+213 + Homo sapiens	
+0.00 +	
-0003 - Oyolin H ribozyme binding site	
<400 ≥ 0901	
coatittate tagtgeete	19
ssasterass cagogoses	<b>4</b> '
K210 × 7922	
3211 × 19	
+(212 + DMA	
H213v Homo sapiens	
12137 Rollio Sapiens	
k(210 k	
<pre><!--!!!3 --> Cyplin E ribozyme binding site</pre>	
4400 + 0922	
Attituatota gtgootoca	19
H210 A 2923	
<211 + 19	
<212 > DNA	
<pre></pre>	
.215 Monto Sapiens	
+(2.00 %	
+2033 Cyclin H ribozyme binding site	
- 400 × 2923	
ctaqtgcctc cagggctgg	19
HD19N 2M24	
-0115 19	
HOIDS INA	
+C135 Heme sapiens	
H22.05	
HIII W Cyclin H ribczyme binding site	
1, out in transpired bring bree	
4400% 2924	
ggctggaatt actatggaa	19

H210 + 2925 H211 + 19 H212 + DNA H213 + Homo sapiens	
+220 + +2221 H ribozyme binding site	
-:400% 2%25 gotggaanta otatggaaa	19
+110 + 1926 +211 + 19 +212 + DNA <213 : Homo sapiens	
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k400% 1927 atggaaagtt atttatoag	19
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
82200 -0220- Cyclin H ribozyme binding site	
+40008 2028 tggaaagtta tttatcaga	19
+1108 2929 +1118 19 +1118 DNA +1138 Homo sapiens	
+1200+ +1223+ Cyclin H ribozyme binding site	

<pre>0400 + 2329 gaaagttatt tatcagaga</pre>	19
<pre>%210 &gt; 2930 %211 + 19 %212 + DNA %213 &gt; Homo sapiens</pre>	
<pre><pre><pre><pre><pre></pre></pre><pre><pre><pre><pre><pre></pre></pre><pre><pre><pre><pre><pre><pre><pre>&lt;</pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>&lt;400% 1930 aaayttattt atcagagag</pre>	19
+210 × 2931 +211 × 19 +212 > DNA +213 > Romo sapiens	
+220 + +223 > Cyclin H ribozyme binding site	
H400% 2931 aagttattta toagagagt	19
<pre>CD10+ 1930 CD11&gt; 19 CD12&gt; DNA CD13&gt; Homo sapiens</pre>	
<2200>	
H400+ 2932 qttatttato agagagtot	19
<pre>#2109 0933 #22109 19 #22109 DNA #22130 Homo sapiens</pre>	
+00000 +00030 Cyclin H ribozyme binding site	
+:400:- 1933 tcadagadto tgatgotga	19
+02100+ 0.434 +02110+ 19 +02110+ DNA	

-:213 · Homo sapiens	
< 220 +	
<pre><!--233 · Tyclin H ribozyme binding site</pre--></pre>	
aadaqaabtt gootgtcac	19
<210 + 1935	
S211 + 19	
-0.212 + DNA	
-CC13 - Homo sapiens	
+:200 +	
<2003 - Cyclin H ribozyme binding site	
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ottgootgto acagttact	19
k210 + 2936	
K211 + 19	
<212 - DNA	
<213 · Homo sapiens	
<2205	
<2223 - Oyclin H ribozyme binding site	
<400 \ 2936	
tytoacaytt actagatat	19
cycoacayco actagatac	1.3
K210 + 2937	
K211 × 19	
SOIDS DNA	
<pre>&lt;213 - Homo sapiens</pre>	
K220%	
R223 - Tyclin H ribozyme binding site	
-:400% 2937	
qtdadagtta otagatata	19
+:::10:> ::2938	
+:211% 19	
-12121- ONA	
+12130 Homo sapiens	
+: <u>2.2</u> 0:-	
H2270 Cyclin H ribozyme binding site	
Nazzur Wyerin n ribozyme binding site	
4400 × 2938	
acagttacta gatataatg	19

+0010 + 0039 +0010 + 19 +0010 + DNA +0013 + Homo sapiens	
+0020 + +0023 + Oyolin H ribozyme binding site	
0400 - 2939 ttastagata taatgaaaa	19
+0210 + 2940 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
<2200 <223> Cyclin H ribozyme binding site	
H400 - 2940 actagatata atgaaaagc	19
<pre>\$210 \cdot 2941 \$211 \cdot 19 \$211 \cdot 8 DNA \$2013 \cdot Home sapiens</pre>	
<pre>+DD0s &gt;DD3s Cyclin H ribozyme binding site</pre>	
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SC10+ 2942 SC11+ 19 SC112+ DNA SC13- Home sapiens	
+:220:- +:223:- Cyclin H ribozyme binding site	
k4000-2942 dagawantta gtaaagaag	19
+2100+ 2943 +0110+ 19 +0100+ DNA +0130+ Homo sapiens	
+12/2B*>	

-:223 - Cyplin H ribozyme binding site	
กสาย การและ 2943 ลิลลองปลัฐบิล aagaagtat	19
<pre>\$210 \ 2044 &lt;221 \ 10 &lt;211 \ ONA &lt;213 \ Homo sapiens</pre>	
<pre>schips <ild> <ignormalize{children} <="" pre=""> <pre>children</pre> <pre>ste</pre> </ignormalize{children}></ild></pre>	
k400% 2944 taaaqaaqta tgaaccacc	19
<pre>&lt;010 * 0945 &lt;011 &gt; 19 &lt;010 &gt; DNA &lt;013 &gt; Homo sapiens</pre>	
<pre>&lt;220s &lt;223s Oyolin H ribozyme binding site</pre>	
<pre>&lt;400&gt; 2945 cacccagate tgaagaagt</pre>	19
<pre>&lt;210 * 1946 &lt;211 * 19 &lt;212 * DNA &lt;213 * Homo sapiens</pre>	
klldds klldds Gydlin H ribozyme binding site	
k4008 [946] tawaqaagtt gotgttotg	19
%2100-2947 *2110-19 %2100-DNA *22130-Homo sapiens	
+(210)+ +(210)+ Cyclin H ribozyme binding site	
H4000- 1447 agtiqonqtu otgaaacag	19
+(210)+ 2948 +(211)= 19	

-:212 - DNA	
Hill3 · Homo sapiens	
+1120 +	
-0.023 · Cyclin H ribozyme binding site	
+:400 + 2943	
gttgbtgttc tgaaacaga	19
+210 × 2349	
±211 × 19	
HALLON DNA	
-MAIR Homo sapiens	
<pre>%ilit@x %ilit3% Cyclin H ribozyme binding site</pre>	
) - Cyclin in Fibozyme binding Site	
H400> 2949	
aacaqaaqtt ggagcgatg	19
<210 > 2950	
<211> 19	
KULIA ENA	
<213> Homo sapiens	
ALTON HUMO Sapiens	
editités	
SCACO Tyclin E ribozyme binding site	
×400× 2950	
gagogatoto attotgotg	19
K2108 0951	
K2115-19	
S212 + ENA	
KLIBA Homo sapiens	
+(ML)0.+	
SCONS Cyclin H ribozyme binding site	
(400) - 1,61	
caatqtcatt ctgctgage	19
Jun 1901, Will Gogolyayo	19
+2108 2952	
H211H 19	
HILLER DNA	
+113+ Homo sapiens	
(200)	
+2%%+ Cyclin H ribozyme binding site	
*.4UU2 - 2. *DZ	

gatgtcattc tgctgagct	19
<pre>#010 + 2953 #011 + 19 #0105 DNA #013 Homo sapiens</pre>	
+320 + +323 + Syclin H ribozyme binding site	
राव00 + 0953 tgotgagett goacttaac	19
<pre>&lt;010&gt; 0354 &lt;011&gt; 19 &lt;012&gt; DNA &lt;013&gt; Homo sapiens</pre>	
<2200 <223% Cyclin H ribozyme binding site	
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<2108 0955 <2118 19 <2128 DNA <2138 Homo sapiens	
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R4008 2955 ottgeactta aegtaatea	19
<pre>&lt;210&gt; 2956 &lt;2118 19 &lt;2128 DNA &lt;2138 Homo sapiens</pre>	
<pre><pre><pre><pre><pre><pre></pre></pre><pre><pre><pre>CCCSS</pre></pre><pre>Cyclin H ribozyme binding site</pre></pre></pre></pre></pre></pre>	
<pre><id400:- 1:356="" acttaacata="" atcacgaag<="" pre=""></id400:-></pre>	19
+00100+1957 +0010+19 +0210+100A +0210+Hemo sapiens	

<pre>HARD* Hardward</pre>	
H400 + 2957 taabgtaatb acgaagaag	19
H210 + 2958 H211 + 19 H212 + DNA H213 + Hamo sapiens	
<pre>-0.200.03 - Cyclin H ribozyme binding site</pre>	
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<210 ~ 2959 <211 - 19 <310 > DNA <313 - Homo sapiens	
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+0010 + 0960 +0011 + 19 +0010 > DNA +0013 > Homo sapiens	
K2200 K2218 Cyclin H ribozyme binding site	
-14001- 1960 atqatgatta ogtotoaaa	19
+00160+ 0961 +00110+ 19 +00120+ DNA +00150+ Homo sapiens	
+.2260+ +.2230+ Cyclin H ribozyme binding site	
-04000- 2361 tqattaoqto toaaagaaa	19
+(210)2962	

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.:211 - 19
 3212 - DNA
+1213 > Homo sapiens
<2200>
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-:4005 2962
attacgtoto aaagaaato
                                                                         19
<210 → 2963
<3.11 \times 19
⊴012> 5NA
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                                                                         19
caaagaaato caaacatga
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                                                                         19
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< 112 > DNA
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<22205
<2235 Cyclin Al ribozyme binding site</pre>
-4005 2965
                                                                        19
ggtgttgttc cggacacat
RD105 0966
·#115 19
-if 120 - DNA
+0.013 Homo sapiens
·22205
+3223> Cyclin Al ribozyme binding site
```

•

-:400 - 0966 oggađasata gaaagataa	19
+010 + 0367 +011 + 18 +010 + DNA +013 * Homo sapiens	
<pre><did0> <id0)> Gyolin Al ribozyme binding site</id0)></did0></pre>	
იქინა _ 1967 tagaaagata acgacggga	19
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04000-0968 gagaacatt tggggtaa	19
+2108 2969 +2218 19 +2212 = ENA	
<pre>&lt;213&gt; Homo sapiens</pre>	
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-:400:- 1969 agasacantt ggggtocag	19
H2168 0970 H2118 19 H2108 DNA H2138 Homo sapiens	
-02205	
-CLICES Cyclin Al ribozyme binding site -C4000-L470	
qttaggate caggcaggt	19
+00100+ 1471 +00110+ 14 +02100+ DNA +02132 Home saniens	
1.7 1.72 NOTE: 5dD1E15	

	- x220%	
	<pre><!--223 * Typlin Al ribozyme binding site</pre--></pre>	
	<400 ≤ 2971	
	osggoaggtt ttgggggoot	19
	<210 × 2972	
	K211x 19	
	<pre><c12> DNA</c12></pre>	
	<pre><i213> Homo sapiens</i213></pre>	
	K0208	
	<pre>&lt;1.23&gt; Cyclin A1 ribozyme binding site</pre>	
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	<210> 3973	
	<211> 19	
	<212> DNA	
	<213> Homo sapiens	
	<220>	
14 1 21 41	<223> Cyclin Al ribozyme binding site	
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	<211> 19	
-	<212> DNA	
	<2135 Homo sapiens	
	<:220×	
	<pre>&lt;223   Cyclin A1 ribozyme binding site</pre>	
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	ttggggeete etgtetggt	19
	+2108 J975	
	+2118 (19)	
	K2125 JNA	
	(C15) Homo sapiens	
	+(220)+	
	<pre>&lt;22.38 Oyclin Al ribozyme binding site</pre>	
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	geeteetgte tggtgggag	19

<pre>%210 * 2976 %211 * 19 %212 * DMA %213 * Homo sapiens</pre>	
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K1805 K1885 Gydlin Al ribozyme binding site	
k4005 0979 agadoggott tocogoaat	19
<pre>%310</pre>	
02230 Cyclin Al ribozyme binding site	

√400 > 2380	
quedagetti edegouate	19
+:210 + 1981	
4211 • 19	
→1011 + DMA	
+213 + Homo sapiens	
+(220)+	
<223 - Cyclin A1 ribozyme binding site	
4400 + 3981	
accggctttc ccgcaatca	19
K210 X 2982	
K2118-19	
<212 > BMA	
<pre><dib> Home sapiens</dib></pre>	
K220 -	
<pre>&lt;223&gt; Cyclin Al ribozyme binding site</pre>	
K400N 2982	
topogoaato atgtacoot	19
<210 + 2983	
<211> 19	
+1212 + INA	
SC13 - Homo sapiens	
<0.00 \$	
<pre><dd3> Cyclin Al ribozyme binding site</dd3></pre>	
+.400.5 (1933)	
caatcatgta cootggato	19
NO10% 2 484	
HOLLS 19	
HOLLIN INA	
- MII) Home sapiens	
+:010ts	
+CCC3+ Cyclin Al ribozyme binding site	
-:400:- 1984	
accongrate tittatigg	19
-M10: 1985	
H22117- 19	
21 "> 5NA	

-:213 → Homo sapiens	
<pre>-1220&gt; -1223&gt; Gyclin Al ribozyme binding site</pre>	
<pre>-(400&gt; 1985 cotggatott ttattgggg</pre>	19
<pre>&lt; 010 &gt; 0986 &lt;011 &gt; 19 &lt;010 &gt; DNA &lt;013 &gt; Homo sapiens</pre>	
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<210> 2987 <211> 19 <212> DNA <213> Homo sapiens	
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<210> 2988 <211> 19 <312> PNA <313> Homo sapiens	
<2200> <2203> Cyclin Al ribozyme binding site	
<400% 2988 mgatottota tigggggot	19
<pre>%210&gt; 2989 %211&gt; 19 %2128 DNA %2138 Homo sapiens</pre>	
-00100- -0223: Cyclin Al ribozyme binding site	
+:400% 2989 atcttttatt gggggctgg	19

H210 + 2990 H211 + 19 H210 + DNA H213 + Homp sapiens	
<pre><l20. *="" <c223="" al="" binding="" pre="" ribozyme="" site<="" typlin=""></l20.></pre>	
c400 × 2330 qayaagaqta totoagotg	19
+0010 > 0991 +0011 > 19 +0012 > DNA +0013 > Homo sapiens	
<220> -(223> Cyclin Al ribozyme binding site	
√400 → 2991  gaagagtato teagetggg	19
<pre>0210 &gt; 2992 0211 &gt; 19 0212 &gt; DNA 0213 &gt; Home sapiens</pre>	
+22.05 +22.35 Cyclin Al ribozyme binding site	
k400% 2992 agagtatoto agotgggaa	19
+0210 × 1993 +0211 × 19 +0012 × DNA +0013 × Home sapiens	
+0000+ +0003+ Cyclin Al ribozyme binding site	
+:400:+ 3993 accoggodto ccagattto	19
+0010+ 0994 +0010+ 14 +0010+ DNA +0013+ Homo sapiens	
~22A5	

HRRR - Cyclin Al ribozyme binding site	
H400+, 994 Stabbagatt togtottoc	19
北210 - 1995 北211 - 19 北211 - BMA	
+213 + Homo sapiens	
+120 + +223 + Cyclin Al ribozyme binding site	
<pre>(400 + 1995) toccagattt cgtottoca</pre>	19
<210 - 2996 <211 - 19	
<pre><c10 +="" <c13="" dna="" homo="" pre="" sapiens<=""></c10></pre>	
<220% <223 - Cyclin Al ribozyme binding site	
k400 - 3996 popagatito gioticoag	19
K210 + 1997 K211 + 19	
<pre>%C12 + DNA %C13 + Home sapiens</pre>	
<pre>&lt;2008 &lt;2008 &lt;2003- Cyclin Al ribozyme binding site</pre>	
-:400:- 2997 adatitodin tiqoaqoaq	19
8210% 1998	13'
HULLY 19 HULLY DNA	
SULTED Homo sapiens	
+0.00+ +0.00+ Cyclin Al ribozyme binding site	
+:400+ 2998	
attingtott deageagea	19
<pre>&lt;210&gt; 2393</pre>	
+:211> 13	

HC212 + DNA HC313 + Homo sapiens	
+:220 + +:220 - 1   1   2   2   2   2   2   2   2   2	
-:400 - 2999 titiogtotic cagcagoag	19
+1010 × 3000 +1011 × 19	
+2012 + DNA +2013 + Homo sapiens	
<pre>&lt;020&gt; &lt;113&gt; Cyclin Al ribozyme binding site</pre>	
-:400> 3000 cogtqqaqtc tgaagcaat	19
<pre></pre>	
<pre><pre><ili><pre><ipre></ipre></pre></ili></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre>	
<pre>&lt;:2208 &lt;:2238 Cyclin Al ribozyme binding site</pre>	
0400> 3001 gagtggagtt gtgctggct	19
<pre>       <br <="" td=""/><td></td></pre>	
+212> FNA +213> Homo sapiens	
+:223+ +:223+ Cyclin Al ribozyme binding site	
04008-3002 gtgotggota cagtggccc	19
+210+ 3003 +211+ 19 +210+ FNA	
<pre>%2130 Hamo sapiens</pre>	
<pre>&lt;1200+ &lt;2200+ Cyplin Al ribozyme binding site</pre>	
(400): 3303	

gooogaggto oogatgott	19
+:210 + 3004 -:211 + 19 -:212 + DNA	
HOME FOR Sapiens	
<pre>SCORT</pre>	
+:400 + 3004 -cocyatystt gtcagatac	19
H2104 3005 H211> 19 H211> DNA	
<pre>%Nlb&gt; Homo sapiens</pre>	
<pre>&lt;1.10&gt; &lt;223&gt; Cyclin Al riboxyme binding site</pre>	
<400> 3005 gatgottgto agatactoa	19
<010> 3006 +0015 19 <010> DMA	
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<pre><pre><pre>&lt;223&gt; Cyclin A1 ribozyme binding site</pre></pre></pre>	
<400% 3006 ttgtbagata otbaccaga	19
<2108 3007 80118 19 80108 DMA	
+0138 Homo sapiens	
+00008 +0003+ Cyclin Al ribozyme binding site	
+:4005 3607 *dagatanto accagagos	19
+.216 % 3008 +:211:- 13	
HIII DNA H2I3> Homo sapiens	

-(22))	
+:223 + Cyclin A1 ribozyme binding site	
-:400 - 3008	
ggobaggato coccgoaga	19
HIZ10 + 3009	
*1211 × 13	
HOTO: DNA	
+0.13 - Homo sapiens	
+:0.20 +	
+MM3+ Cyclin Al ribozyme binding site	
<400 + 3009	
gacantgota gggotgota	19
<0.10 + 3010	
KULLA 19	
<pre><c10 -="" dma<="" pre=""></c10></pre>	
K213 - Homo sapiens	
Letter Reduce Saprens	
<2200 +	
<pre>&lt;123 · Cyclin Al ribozyme binding site</pre>	
<ul><li>&lt;400 ≥ 3010</li></ul>	
agggotgota actgcaaat	19
+:210 + 3011	
H211 + 19	
KC12 + DNA	
<pre>sc. 13&gt; Homo sapiens</pre>	
CONTROL OF THE RESIDENCE OF THE STATE OF THE	
-CMC3- Cyclin Al ribozyme binding site	
x(400 × 3011	
atoggoagta caggaggac	19
-010 m - 3012	
-II 1 II - 19	
HUIDALIN DNA	
40130 Home sapiens	
1927.Ar.	
12223 Cyclin Al ribozyme binding site	
+(40):>=3012	
ccaddggate acaagaate	19
+21m2 3n13	

K.::11 > 19	
-1.112 · DNA	
-1.113 · Homo sapiens	
<u.20 +<="" p=""></u.20>	
<pre></pre>	
-(400 + 3013	
oacaagaato aggtgttat	19
<210 + 3014	
K211 × 19	
+213 + 5MA	
-213 - Romo sapiens	
KU209	
<223 Cyclin Al ribozyme binding site	
<400 ± 3014	
atcaqqtqtt attctggat	19
-:210>-3015	
<211 ← 19	
KO10 + DNA	
<013 Homo sapiens	
K120 +	
SDD3 - Cyclin Al ribozyme binding site	
<400% 3015	
traggtgtta ttotggato	19
:2180-3016	
SM11:- 19	
HALLO DNA	
CD130 Hemo sapiens	
- (mp.m.)	
<pre>:IIII30 Cyclin Al ribozyme binding site</pre>	
(40c)- 3016	
aqqtqttätt otggatoag	19
02105 3017	
01110-19	
COLD DNA	
ALLE: Homo sapiens	
kares	
<223> Cyclin Al ribozyme binding site	

-(400 - 3017 ggtgttatto tggatcaga	19
+0010 + 3013 +0011 + 19 +0012 + DNA +0013 + Homb sapiens	
-0.20	
<pre><i400 -="" 3018="" agaaaatgo<="" attotggato="" pre=""></i400></pre>	19
+:210 + 3019 +:211 + 19 -:210 + DNA	
<pre>#213 Homo sapiens</pre>	
<pre>&lt;0200&gt; <pre><pre><pre><pre>&lt;0.23 - Cyclin A1 ribozyme binding site</pre></pre></pre></pre></pre>	
-:400 > 3019 aaaatgoott cootooago	19
+210 > 3020 +211 + 19	
+:212 + DNA +:213 + Homo sapiens	
+2200.4 <223% Cyclin Al ribozyme binding site	
+:400:+ 3000 awatgootto cotocaget	19
+(210)+ 3001 +(210)+ 19	
HIMTO DMA HIMTO Home sapiens	
+00000 +00035 Cyclin Al ribozyme binding site	
+400:-30::1 qcc::ccctc cagctggaa	19
+0010 + 3002 +0010 + 19 +0010 + DNA	
H213P Homo sapiens	

<pre>&lt;:220 * &lt;:223 * Gyplin Al ribozyme binding site</pre>	
+: <b>4</b> 00+ 3022	
gaaageasto eetgaetgt	19
<:2108-3123	
-0.11 × 19	
FIG 12 × DNA	
+2.13 m Homo sapiens	
+12.00 ×	
4223 % Gyplin Al ribozyme binding site	
<400> 3023	
atgtqgqqta caagagcc	19
+1210> 3024	
<211> 19	
COLLS DUA	
+213> Homo sapiens	
<pre>&lt;223&gt; Cyclin Al ribozyme binding site</pre>	
+:400 × 3024	
agcaagggtt tgacatcta	19
+::10 + 3025	
+011+ 19	
SDID> ENA	
$ ilde{c}$ 213 $ ilde{c}$ Homo sapiens	
+1020s	
-223 Cyclin Al ribozyme binding site	
-:4005-3005	
qcaaqggttt gacatctac	19
+0010N 30D6	
+:::110 19	
+:D12:N	
and the transportation of the transport	
6.02 (b)	
H213H Cyclin Al ribozyme binding site	
-:400:- 3026	
gtttgacatc tacatggat	19

<pre><d210 +="" 19="" 3027="" <d211="" <d213="" <d312="" dna="" homo="" pre="" sapiens<=""></d210></pre>	
+0000+ +0003+ Gyalin Al ribozyme binding site	
0400 + 3027 tigapatota datggatga	19
<pre>&lt;:210 + 3028 &lt;:211 + 19 &lt;:212 + DNA &lt;:213 &gt; Homo sapiens</pre>	
<2230 Cyclin Al ribozyme binding site	
<400> 3028 ggatgaacta gagcagggg	19
<210× 3029 <211× 19 <212× DNA <213× Homo sapiens	
-Cus. -Cus. Cyclin Al ribozyme binding site	
<pre>+:400 x :3029 acagetgete ggtcagaga</pre>	19
+2108 3030 +2118 19 +2118 DNA +213 Homo sapiens	
+2rus +27rus Cyclin Al ribozyme binding site	
-:400% 3030 	19
+:210:- 3031 +:211:- 19 +:21:- DNA +:21:- Homo sapiens	
<pre>%2200: %223% Cyclin Al ribozyme binding site</pre>	

<pre>-:400 + 3031 pyatygcatt tgaggatgt</pre>	19
+010 + 3030 +011 + 10 +010 + DNA +0210 + Homo sapiens	
<pre><d200x <d220x="" <d223=""> Oyolin Al ribozyme binding site</d200x></pre>	
-:400 = 3032 gatogcatt: gaggatgtg	19
<pre>&lt;010&gt; 3033 &lt;011&gt; 19 &lt;010&gt; ONA</pre>	
<pre>%113&gt; Homo sapiens</pre>	
<220> <225> Cyclin Al ribozyme binding site	
<400> 3033 aqqatgtqta tgaagtaga	19
<210 > 3034 <211 > 19 <212 > DMA	
<pre>&lt;213&gt; Homo sapiens</pre>	
<2208 <2238 Cyclin Al ribozyme binding site	
+1400% 3034	19
atatqaaqta gacacoggo	13
<pre>&lt;210&gt; 3035 +211&gt; 19 +210&gt; DNA</pre>	
HEILD DUA HEILD Homo sapiens	
<22.08	
+CDCC+ Cyclin Al ribozyme binding site	
+14005 3035	19
oggdacantd aagtoagac	19
+0.11(a) 3036	
+00110+ 19 +00120+ DNA	

-:213 · Homo sapiens	
+12.20 +	
<pre>*M.M3 * Cyclin Al ribozyme binding site</pre>	
√400 ← 3036	
eastsaadts agacetgca	19
+210 + 3637	
+211 + 19	
K212 + DNA	
<pre>c213 - Homo sapiens</pre>	
< 2.1.0 +	
+213 + Cyclin Al ribozyme binding site	
<400 > 3037	1.0
abbtqcabtt obtgbtgga	19
<210 × 3038	
<211 - 19	
RC1C> DNA	
<213> Homo sapiens	
K220 ×	
+223 + Gyolin Al riboxyme binding site	
<400 ≤ 3038	
ectquactto otgotggat	19
<210 > 3039	
<011 × 19	
<pre><pre></pre></pre>	
<pre>*Dl3* Homo sapiens</pre>	
kaza -	
<pre><pre></pre>Clist Cyclin Al ribozyme binding site</pre>	
840CH 3033	
etgetggatt teaacacag	19
+2100-3040	
k0115-19	
+12.170 + 12MA	
+Illy - Homo sapiens	
+20c)+	
K2000 Cyclin Al ribozyme binding site	
tgctggattt caacacagt	19

HC10+ 3041 HC11+ 19 HC10+ DNA HC13+ HCMC sapiens	
<pre>&lt;220 &gt; <pre></pre> <pre>&lt;2220 *</pre> <pre>&lt;2223 * Cyclin A1 ribozyme binding site</pre></pre>	
<pre>&lt;4008 3041 gotggattto aacacagtt</pre>	19
<pre>&lt;310&gt; 3042 &lt;311&gt; 19 &lt;312&gt; DNA &lt;213&gt; Homo sapiens</pre>	
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<210> 3043 <211> 19 <210> DNA <213> Homo sapiens	
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<pre>K0108 3044 K0118 19 K0118 EMA K0138 Homo sapiens</pre>	
<pre>&lt;2200&gt; &lt;2003 Cyclin Al ribozyme binding site</pre>	
k4008 3044 adadagttto doctatget	19
+00100+ 3045 +0010+ 104 +0010+ DNA +0010+ Homo sapiens	
+12121Qt>	

HRRRY Cyclin Al ribozyme binding site	
+:400 + 3045	
gttf poosta tgotggtag	19
√310 - 3046	
si211 + 19	
<pre>%210 + DNA</pre>	
8213 - Homo sapiens	
+12.00 +	
<pre>&lt;223 + Cyclin A1 ribozyme binding site</pre>	
₹400 + 3046	
tatwotggta gattoatot	19
<210 > 3047	
<211 × 19	
<210 > DNA	
<213> Homo sapiens	
<2005	
<pre>&lt;2.13&gt; Cyclin A1 ribozyme binding site</pre>	
<400> 3047	
ctqqtagatt catctctcc	19
KD108-3048	
<2118-19	
K2128 DNA	
<pre>&lt;2135 Homo sapiens</pre>	
<2.00%	
<pre>%ABSO Cyclin Al ribozyme binding site</pre>	
k(400) - 3048	
tggtagatto atototoot	19
8/210/- 3/49	
-00110-19	
AMD SMA	
<pre>SOLIC Homo sapiens</pre>	
-conde-	
-Chias Cyclin Al ribozyme binding site	
+14000+ 3049	
tagarteate tetectete	19
+II:10:- 3050	
+211× 19	

HOME Sapiens	
<pre><dincelled +="" al="" binding="" carolin="" pre="" ribbzyme="" site<=""></dincelled></pre>	
<pre>4400 + 3050 gatteatete tectetees</pre>	19
#210 + 3051 #211 + 19 <212 + DNA #213 + Homo sapiens	
<220> <223 - Cyclin A1 ribozyme binding site	
<400> 3051 tteatetete eteteceag	19
<210> 3052 <211> 19 <212> DNA <213> Homo sapiens	
<2205 <2235 Cyclin Al ribosyme binding site	
<400% 3052 atotototo toccagtot	19
<210% 3053 <211% 19 <210% DNA <213. Homo sapiens	
<pre>&lt;2208 &lt;2238 Cyclin A1 ribozyme binding site</pre>	
k400s 3053 dictiocidic coagiciga	19
<pre><c108 +00108="" +00118="" 19="" 3054="" <c108="" dna="" homo="" pre="" sapiens<=""></c108></pre>	
<pre>+DENCO: +DENCO: +</pre>	
÷(400)÷ 3054	

.

totoocagto tgaagatat	19
<pre>&lt;210 &gt; 3055</pre> <pre>&lt;211 + 19</pre>	
CD1D+ DNA CD13+ Homo sapiens	
<pre>&lt;2200* &lt;223* Oyolin Al ribozyme binding site</pre>	
-400 / 3055 totquagata tatocagto	19
<pre>&lt;210 + 3056 &lt;211 + 19 &lt;212 &gt; DNA</pre>	
+213 + Homo sapiens	
-:220> -:23:3- Cyclin Al ribozyme binding site	
<400> 3056 tgaaqatata tocagtott	19
<210 × 3057 <211 × 19	
HORA Homo sapiens	
<pre><din0 +="" <idin3=""> Cyclin Al ribozyme binding site</din0></pre>	
<400° 3057 aagatatato cagtottgg	19
HIZ 10 8 (E058) HIZ 11 + 14	
HOLDY ENA HOLDY Homo sapiens	
<pre>&lt;:220 - &lt;:223&gt; Cyclin Al ribozyme binding site</pre>	
-:40c5 3058 atatocaqto tiggoacag	19
-0.160-3059 -0.110-14	
42115 DNA	
K2130 Homo sapiens	

+:22a +	
+:223 + Cyclin Al ribozyme binding site	
₹4±0 + 3±59	
atobagtett ggcacagat	19
+210 + 3060	
8211 - 19	
H212 + DMA	
<pre>si213 - Homo sapiens</pre>	
kunia -	
<pre><!--d.3 + Cyclin Al ribozyme binding site</pre--></pre>	
<400 ≠ 3560	
agatutgata aatgtgact	19
<210: 3061	
<211> 19 <2125 DNA	
RCHAR BOMA RCHAR Homo sapiens	
ALTO FIGURE Sapiens	
\$0.00 S	
<pre>&lt;.3.23   Cyclin Al ribozyme binding site</pre>	
s(400 × 3061	
tgactgaata tgctgaaga	19
+0.10 % 3062	
K011 + 19	
HARLES DNA	
<22135 Home sapiens	
k2200k	
<pre>&lt;223&gt; Cyclin Al ribozyme binding site</pre>	
-:400:> 3052	
tgaagaaatt tatcagtac	19
+:::10x 3063	
-C110-19	
+M1M+ DNA +M13% Homo sapiens	
Salves nome sapiens	
(III.20)s	
0.230 Cyclin Al ribozyme binding site	
qaadaaattt atcagtacc	19
405 tipe - 3064	

.

21.0 Falm	
COLO - DNA	
-(213 - Homo sapiens	
₹220 ×	
-CD23 - Cyplin Al ribozyme binding site	
-(400 + 3064	
aaqaaattta toagtaoot	19
-(216 + 3065)	
H211 + 19	
-0212 + DNA	
$\leq 2.13 imes$ Homo sapiens	
*:220 ·	
+223 + Cyclin Al ribozyme binding site	
84400 + 30 <b>65</b>	1.0
vaaatttato agtacotta	19
+:210 + 3066	
KD11 + 19	
HOTO HIDNA	
+C213 · Homo sapiens	
+12.11.0 +	
+33.3 + Cyclin Al ribozyme binding site	
<400 + 3066	
tttatoagta oottaggga	19
H210H 3067	
K211 + 19	
COLOR DNA	
-C2130 Homo sapiens	
H0000	
HIGHE Cyclin Al ribozyme binding site	
-04000- 3067	
toadtacett agggaaget	19
HIDDON 3068	
RI2010-19	
HODDH DNA	
Homo sapiens	
H22016	
<223> Cyclin Al ribozyme binding site	

<pre>&lt;4400 * 3063 bagtacotta gggaagetg</pre>	19
<pre>#010 + 3069 #011 + 19 #012 + DNA #013 + Homo sapiens</pre>	
+(120 + + Cyclin Al ribozyme binding site	
ाद्मण्याः । ३०६५ ayotgaaata   aggcacaga	19
<210 + 3070 <211 > 19 <212 > DNA	
<2213> Home sapiens	
<pre>&lt;223&gt; Cyclin A1 ribozyme binding site &lt;400&gt; 3070</pre>	10
aagcacacta catgaagaa <210> 3071	19
+211> 19 +212> ENA +213> Humo sapiens	
<pre>&lt;0.00&gt; &lt;0.03&gt; Cyclin Al ribozyme binding site</pre>	
(4008-3071 godagadato acggaaggo	19
#2108 3072 #211> 19 #2118 DNA #2138 Homo sapiens	
-0000- -0003- Cyclin Al ribozyme binding site	
e400: 307: gogcacgatt otggtggac	19
#216: 3073 #210: 19 #210: DNA #215: Home sapiens	

	:220 %	
	<pre>&lt;223 * Oyolin Al ribozyme binding site</pre>	
	+400 + 3073	
	equality tggtggact	19
	<pre><pre><pre></pre></pre></pre>	
	-(211 + 19	
	+210 → DNA	
	<pre>#213 * Homo sapiens</pre>	
	K220 +	
	<223 - Syclin Al ribozyme binding site	
	<400 + 3074	
	gytqdagatt ggggaagaa	19
	<210 × 3075	
	<0.11 + 19	
	KO12 + DNA	
	(213) Homo sapiens	
4	4.000 ×	
*** *** **	+333 + Cyclin Al ribozyme binding site	
-	-:400 + 3075	
	qqqaagaata taaacttog	19
	+210.+ 3076	
	<2115 19 (Control of the Control of	
	K2125 DNA	
÷.	-C2130 Humo sapiens	
	+(MD6)+	
	<pre>CD230 Cyclin A1 ribozyme binding site</pre>	
	-:400:- 3076	
	qaadaatata aacttogag	19
	H2160 3077	
	(2115-19)	
	KO120 DNA	
	-CM130 Romo sapiens	
	kmm to k	
	-CECEN- Cyclin Al ribozyme binding site	
	<400> 3077	
	atataaactt cgagcagag	19

<pre><d210 +="" 19="" 3078="" <d211="" <d212="" <d213="" dna="" homo="" pre="" sapiens<=""></d210></pre>	
<pre>&lt;:220 + &lt;:223 + Gyolin Al ribozyme binding site</pre>	
<pre>+(400 + 3078) tataaactto gagcagaga</pre>	19
+010 + 3079 +011 + 19 +011 + DNA +013 > Homo sapiens	
<pre><dio> <doo> <doo <="" pre=""></doo></doo></dio></pre>	
#400% 3079 agaddotgta totggotgt	19
<pre>&lt;010 &gt; 3080 &lt;011 &gt; 19 &lt;010 &gt; DNA &lt;013 &gt; Home sapiens</pre>	
<22008 <223% Cyclin Al ribozyme binding site	
k400% 3080 addotgtato tggotgtoa	19
<pre>%1108 3081 %.118 19 %.178 DNA %.130 Homo sapiens</pre>	
-C220- -C238 Cyclin Al ribozyme binding site	
-4000-3081 totamotata aacttoota	19
+0110+ 3:82 +0210+ 1+ +0210+ DNA +0210+ Homo sapiens	
<pre><pre><pre>&lt;225&gt; Cyclin A1 ribezyme binding site</pre></pre></pre>	

+:40m + 3082	
stytoaastt cotggacag	19
8210 + 3983	
<211 + 14	
DMA	
+1213 + Homo sapiens	
+MM3 + Cyclin Al ribozyme binding site	
+3490 + 3083	
tgtcaactto otggacagg	19
<210 × 3084	
K2115-19	
KO12 KIDNA	
<pre>&lt;213&gt; Homo sapiens</pre>	
•	
+0.00x	
-1113> Cyclin Al ribozyme binding site	
,	
<400 + 3084	
tqqacagqtt cctttcatg	19
×11.10 × 3085	
K211 + 19	
8010 × 50A	
+1213+ Homo sapiens	
•	
<2208	
<pre>&lt;2235 Cyclin Al ribozyme binding site</pre>	
+400 + 3085	
agadaggtto otttdatgt	19
<2108-3086	
-C118-19	
HILLS ENA	
+2180 Homo sapiens	
+:2005	
<pre><c130 al="" binding="" cyclin="" pre="" ribozyme="" site<=""></c130></pre>	
+(4g0)+-3086	
caggitecti teatgtatg	19
-C.10:- 3087	
0.111:- 19	
HZ12F DNA	

-213 · Homo sapiens	
+1120 +	
-CMD3 - Cyclin Al ribozyme binding site	
+:400 + 3087	
aggitteettt eatgitatgi	19
+210 + 3983	
<211 × 19	
HD1D KIDNA	
<pre>&lt;113 Homo sapiens</pre>	
-:210 /	
<pre><di23> Cyclin A1 ribozyme binding site</di23></pre>	
-:4005 3086	
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·:220>	
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H210× 3090	
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KO1U> DNA	
C135 Homo sapiens	
<0.00°	
CDDA+ Cyclin Al ribozyme binding site	
-040008 30 <b>30</b>	
datutatqto tgttotgag	19
00100 30 <b>9</b> 1	
SD118-19	
CONTROL DNA	
(Clib) Homo sapiens	
and on-	
42000 Cyclin Al ribozyme binding site	
<400× 3291	
tatgtotgtt otgagaggg	19

```
<210 + 3092
\pm 1211 \pm 19
H212 - DNA
%2130 Home sapiens
-1220 ·
+223+ Nolin Al ribozyme binding site
H400> 3090
atgictitte tgagaggga
                                                                          19
+0.10 → 309±
HB11> 19
HARLEY DNA
+2213> Homo sapiens
+ 120>
<223> Cyclin A1 ribozyme binding site
-1400> 3093
                                                                         19
actidoaqotic gtaggaaca
+:210 > 3094
<2217 > 19
3212> DNA
HOME sapiens
-12205
+223> Cyclin A1 ribozyme binding site
<400% 3094
                                                                         19
quaqutoqta ggaacagca
HD 10 + 3095
HI11 - 13
JULIA DNA
+213 · Homo sapiens
·[22:0]·
\oplus 22\, \Im \cdot Cyclin Al ribozyme binding site
-14008-3095
                                                                         19
adagdagdta tgcttttgg
+(21u)+ ±69n
ROLL DNA
< 0.130 \cdot \text{Homo sapiens}
42201-
```

.

<223 · Cyclin A1 ribozyme binding	site
⊴400 + 30 <b>96</b>	
agotatgott ttggottog	19
K210 × 3097	
4211 + 19	
HILL HIDNA	
<2138 Homo sapiens	
<2.20a	
-0003 - Cyclin Al ribozyme binding	site
actatgottt tggcttcga	19
<010× 3098	
<211> 19 <210> DNA	
<213> Homo sapiens	
21.00 Homes Saptems	
<220>	
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<2105 3099 <2115 19	
C212> DNA	
<2138 Homo sapiens	
-	
<2000 >	
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3400.5 3099	
ctittggett egaaatatg	19
C2100- 3100	
0.015 19	
SINI DE DUA	
00135 Homo sapiens	
00260	
い ごご3m Cyclin Al ribozyme binding	site
14001-3100	
''t'qgc'tc gaaatatga	19
:::10:- 3101	
(211): 19	

-::212 ·	DNA	
	Homo sapiens	
	•	
4220-		
	Cyclin Al ribozyme binding site	
	, o	
-(4:1:c+	3101	
		19
22239	c. oguajaga	
-( <u>21</u> 0-	3102	
- 21.		
	Homo sapiens	
	nama oupiono	
<:220 -		
	Cyclin Al ribozyme binding site	
.22.2	tiporin in tracingmo armaing broad	
<400 >	3103	
		19
C 9 3 3 10		-
<2105	3103	
<2.11 -		
<212 ×		
	Homo sapiens	
	Ospi Silo	
<11.10 ×		
	Cyclin Al ribozyme binding site	
	· · · · · · · · · · · · · · · · · · ·	
-14005	3103	
		19
, .		
<210>	3104	
-211:		
-:21.H-		
	Homo sapiens	
	•	
-11:200-		
kin235	Cyclin Al ribezyme binding site	
-14(00)-	3104	
gagat.a	tato otootgaag	19
-12100-	3105	
.:211:-		
	AND.	
	Homo sapiens	
	•	
-1111100-		
.122.35	Cyclin Al ribozyme binding site	
-:400iz	3105	

atatatosts otgaagtag	19
H210 × 3106 H211 · 19 H212 · DNA H213 · Homo sapiens	
+0.00 + +0.000 + Oyolin Al ribozyme binding site	
-:400 × 3100 tootqaaqta gaogagttt	19
<pre>AD10 + 310? AD11 + 19 AD12 + DMA AD13 + Homo sapiens</pre>	
+200 + <2003 - Gyolin Al ribbzyme binding site	
<pre>k400 - 3107 tagacgagtt tgtctatat</pre>	19
<pre>#310 * 3109 #311 * 19 #312 * DNA #313 * Homo sapiens</pre>	
<pre><pre><pre><pre>&lt;220 </pre> <pre>&lt;223 &lt; Cyclin Al ribozyme binding site</pre></pre></pre></pre>	
<pre>&lt;:400 + 3108 agacgagtt: gtctatatc</pre>	19
+1210 + 310 ) +1211 + 19 +1211 + DNA +1213 + Homo sapiens	
+1220 - +1223 - Cyclin Al ribozyme binding site	
+:400:+ 3109 ogagittuto tatatoaco	19
+:21c+ 3:10 +:217+ 14 +:212+ DNA +:213> Home sapiens	

HARRO - HARROS - Cyclin Al ribozyme binding site	
(400 + 311)	
aqtitgtota tatcaccga	19
< 3111	
H211 + 19	
HOID + DNA	
<pre>%213 Homo sapiens</pre>	
Let 5 Name Suprano	
H0.10 N	
RANGE Syclin Al ribozyme binding site	
<400 - 3111	•
tttgtctata tcaccgatg	19
<310 + 3112	
KU11 + 19	
<pre><pre><col/></pre></pre>	
<213 - Home sapiens	
and the state of t	
<220 ←	
<pre>&lt;3.3.3 - Cyclin Al ribozyme binding site</pre>	
<400 × 3112	
tototatato accgatgat	19
-C10+ 3113	
KL112-19	
SCHOOL DNA	
K213> Homo sapiens	
+:2208	
-2230 Cyclin Al ribozyme binding site	
+:400:- 311s	
	19
accdatgata catacacaa	13
H001008 3014	
H2211:- 19	
H011215 DNA	
+22135 Homo sapiens	
+2200	
+2230 Cyclin Al ribozyme binding site	
+(400)- 311;	19
atgatacata cacaaaacg	1.7
+(210)+ 3115	

+0011 + 19 +0012 + DNA +0013 + Homo sapiens	
<pre>HDD0+ HDD3+ Dyclin Al ribozyme binding site</pre>	
H400 + 3115 gabbactitt aasaatgga	19
0210 + 3116 0211 + 19 0212 + DNA 0213 + Homo sapiens	
<pre>&lt;220 - &lt;223 - Cyclin A1 ribozyme binding site</pre>	
k400 - 3116 abaabtgtta aaaatggaa	19
<pre>&lt;210 &gt; 3117 &lt;2211 &gt; 19 &lt;212 + DMA &lt;213 &gt; Homo sapiens</pre>	
<pre>&lt;2200+ &lt;2203+ Cyclin A1 ribozyme binding site</pre>	
k4008 3117 tqqaacaatt gottotgaa	19
<pre>\$210+ 3118 \$211&gt; 19 \$212+ DNA \$2135 Homo sapiens</pre>	
<pre>&lt;1200. <pre></pre> <pre><pre></pre> <pre>Cyclin Al ribozyme binding site</pre></pre></pre>	
-:400:> 3:18 acacttg:tt ctgaaagtt	19
+0.109 3:19 +0.110 1.0 +0.110 DMA +0.130 Homo sapiens	
+:220:- <223:- Cyclin Al ribozyme binding site	

-:400 + 3113 pastigotis igaaagito	19
+:210 + 2100 +:211 × 19 +:212 + DNA +:213 + Homo sapiens	
+12.20 + +12.23 + Cyplin Al ribozyme binding site	
H400 + 3120 totgaaaqtt otagotttt	19
+1210 % 3121 +1211 > 19 +1212 + DMA +1213 % Homo sapiens	
+00008 +1003 + Gyelin Al ribozyme binding site	
+:400 + 3101 otgawagtto tagottttg	19
+010 + 3100 +011 + 19 +011 + DMA +013 + Home sapiens	
<pre>+020 + &lt;223 * Cyclin Al ribozyme binding site</pre>	
нции. 3122 qaaaqttota gottttgat	19
+0010+ 3107 +0017+ 19 +0010+ INA +0013+ Homo sapiens	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
-400-3103 gitolagofi titgatotga	19
+00100+ 3124 +02110+ 14 +00120+ DNA +02130+ Homo sariens	
- Lao- Indus Our action	

<pre><!--220 * *:223 * Cyclin A1 ribozyme binding site</pre--></pre>	
ki400 + 3124 thobagoiit tgatotgao	19
+0210 + 3125 +0311 + 14 +0312 + DMA +0313 + Homo sapiens	
+0000+ +0003+ Gyolin Al ribozyme binding site	
k400% 3125 totagotitt gatotgada	19
<pre>&lt;:210   3106 :2215   19 :2108   DNA :2213   Homo sapiens</pre>	
<pre>kindos kindos cyclin Al ribozyme binding site</pre>	
+:400+ 3126 doi:ttgate tgacagtae	19
<pre>&lt;2108 3127 &lt;211 + 19 &lt;2128 DNA &lt;2138 Homo sapiens</pre>	
RIBON RIBON Cyclin Al ribozyme binding site	
k(4008-3107 totgacagta obaaccacb	19
<pre>#210 + 3123 #2211 + 14 #2212 DNA #2213 Homo sapiens</pre>	
+(220)+ +(223)+ Cyclin Al ribozyme binding site	
-:400> 3123 ccaaccagtt totoottoa	19

<pre>&lt;210 * 3129 </pre>	
HOLLS FOUNA HOLLS Homo sapiens	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
H400+ 312M Gaaccagttt ctccttcag	19
<pre>%210 + 3130 %211 + 19 %213 + DNA %213 + Homp sapiens</pre>	
+:2200 +:223 * Cyclin Al ribozyme binding site	
K400% 3130 aaccagtito toottoagt	19
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<pre>%Clide %Clide %Clide Cyclin Al ribezyme binding site</pre>	
k(400) 3131 ecagtiticis officagias	19
<pre>#0108 3130 #02108 10 #02108 DNA #02130 Home sapiens</pre>	
+22008 +22038 Cyclin Al ribezyme binding site	
+:400:- 313. gtttotoott cagtacttg	19
+0.100 3150 +0.100 19 +0.100 DNA +0.130 Homo sapiens	
:00200- :0223> Cyclin Al ribozyme binding site	

H400+ 3133 tithitcottc agtacttga	19
HD10 + 3134 HD11 + 19 HD12 + DNA HD13 + Homo sapiens	
<pre>%220 * %223 * Gyclin A1 ribozyme binding site</pre>	
-0400 + 3134 + pottoaqta ottgaggog	19
<pre>&lt;210 = 3135 &lt;211 &gt; 19 &lt;212 &gt; DNA &lt;213 &gt; Homo sapiens</pre>	
<pre>&lt;020&gt; &lt;123&gt; Cyclin A1 ribozyme binding site</pre>	
<400> 3135 ttbagtantt gaggogaca	19
\$2108 3136 \$2118 19 \$2128 DNA \$213 - Homo sapiens	
<pre>&lt;2000&gt; &lt;1003 - Cyclin Al ribozyme binding site</pre>	
<pre>&lt;400 &lt; 3136 aqtqtgcqtc aggactgag</pre>	19
<pre>%2100 3137 +02100 19 %. LDS DNA +0.130 Homo sapiens</pre>	
-01008 -01030- Cyclin Al ribozyme binding site	
-:400:- 313" Aachtganta agtacgtag	19
+0.100 3138 +0.11: 19 +0.212: PNA	

+:213 + Homo sapiens	
+12.24 ×	
ACCION Cyclin Al ribozyme binding site	
+4400 + 3133	
tqx:taaqta ogtagosga	19
HU10 + 3139	
H211 × 19	
HILLS DNA	
HD13 + HDmo sapiens	
-0.20 /	
<pre>&lt;123 * Gyolin Al ribozyme binding site</pre>	
<400≯ 3139	
taaqtacqta gosgagotg	19
+210 + 3140	
-211 × 19	
K212 + DNA	
$\pm 213 \pm  ext{Homo}$ sapiens	
+:220 ×	
+:223 - Cyclin Al ribozyme binding site	
-:400 × 3140	
gagotgagto tacttgaag	19
+:210 × 3141	
<211> 19	
COLO DNA	
•213 • Homo sapiens	
kanes	
SCIBS Dyclin Al ribozyme binding site	
<ul><li><a href="https://doi.org/10.1007/j.jps.com/red/">https://doi.org/10.1007/j.jps.com/red/</a></li></ul>	
getgagteta ettgaagea	19
+D100+ 3142	
W2118 19	
HOLDE DNA	
43 139 Hemo sapiens	
H110H	
<pre></pre> <pre>Cyclin Al ribozyme binding site</pre>	
·:400× 5142	
gagtetaett gaageagat	19

HD100 3143 HD11 + 19 HD12 + DNA HD13 + Homo sapiens	
<pre>00.00 color Al ribozyme binding site</pre>	
-:400 + 3143 gaagcagato cattottga	19
+0210 + 3144 +0211 + 19 +0210 + DNA +0213 × Homo sapiens	
<pre>&lt;120 \ &lt;223 - Cyclin A1 ribozyme binding site</pre>	
<pre>0400 - 3144 cagatocatt cttgaaata</pre>	19
<pre>&lt;210 + 3145 &lt;211 * 19 &lt;212 + DMA &lt;213 + Homo sapiens</pre>	
+320 - 4203 - Cyclin Al ribozyme binding site	
<pre>&lt;400&gt; 3145 adatccattc ttgaaatat</pre>	19
+02100+3146 +02110+19 +02100+DMA +02100+Homo sapiens	
-00000- -000000 Cyclin Al ribozyme binding site	
+:400:- 3146 atomattott gaaatatot	19
+01160+ 3147 +01118 14 +0111+ DNA +01130+ Homo sapiens	
+(2.2位)	

4333 Cyclin Al ribozyme binding site	
+:400 + 3147	
totiquata tottootto	19
+C10 + 3148	
42.11 * 19	
HILL DNA	
HUIR - Homo sapiens	
+:0:050 +	
+2003 + Cyclin Al ribozyme binding site	
-:400 + 3148	
tigasatato ticcitcas	19
KO10 × 3149	
<3117-19	
KONDS DNA	
<pre>&lt;113  Homo sapiens</pre>	
<220 +	
<pre>%223 - Cyclin Al ribozyme binding site</pre>	
ALLS - Cyclin Al Tibbzyme binding Site	
<400 ÷ 3149	
quaatatott octtoactg	19
HILLON 3150	
(211) 19	
HD1DH DNA	
K2135 Homo sapiens	
K2008	
<pre><pre><pre></pre>Cyclin Al ribozyme binding site</pre></pre>	
K400X 3150	
aaatatette etteaetga	19
<pre><pre><pre><pre><pre></pre></pre></pre></pre></pre>	
(C11): 19	
COLOR DNA	
CO130 Home sapiens	
.2200	
-2230 Cyclin Al ribozyme binding site	
··4000-3151	
tator toott cactgatag	19
HU100-3152	
+121114 1 1 1	

<pre>4212 + DNA 4213 + Homo sapiens</pre>	
+2.00 + +2.03 + Cyclin Al ribozyme binding site	
<pre>4400 * 3152 atottootto actgatage</pre>	19
+210 + 3153 +211 + 19 +212 + DNA +213 + Homo sapiens	
<pre><pre></pre> <pre><pre><pre></pre> <pre><pre></pre> <pre></pre> <pre><pre></pre> <pre></pre> <pre></pre></pre></pre></pre></pre></pre>	
+:400> 3153 ttbabtgata gotgoagoa	19
+0:10% 3154 +0:11* 19 +0:12* DNA +0:13* Homo sapiens	
<220> <223> Cyclin Al ribozyme binding site	
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<pre>&lt;210 &gt; 3155 &lt;211 &gt; 19 &lt;212 &gt; DNA &lt;213 &gt; Homo sapiens</pre>	
ALING Homo saprens ALING + ALING + Cyclin Al ribczyme binding site	
+400+3155  bagbagett* ttgootggo	19
-210-3156 -211-19 -213- DNA -213- Homo sapiens	
<pre></pre> <pre><pre>&lt;2130 Cyclin Al ribczyme binding site</pre></pre>	
edino. 9156	

agragetttt tgeetggea	19
÷(21) ≥ 3157	
-::11 → 19	
HILL DHA	
+1113 Homo sapiens	
-0000 ·	
-MMB+ Gyalin Al ribozyme binding site	
,	
<400 > 315°	
goaqottttt gootggoaa	19
and the second s	
H210 + 3158	
×2211 × 19	
<212 DNA	
<pre>&lt;213 Home sapiens</pre>	
.215 Holke Sapiens	
K220 /	
<pre>&lt;223    Cyclin A1 ribozyme binding site</pre>	
*225% Cyclin Al Tibozyme binding Site	
K400× 3158	
tqqcaaacta tactgtgaa	19
tygoadacea tactycyda	15
K2108 3159	
K2112-19	
HOLD FONA	
<pre>4213 Home sapiens</pre>	
1213 / Holle Saprens	
<0.20 ×	
*MM3 · Cyclin Al ribozyme binding site	
Cyclin Al Tibozyme binding Site	
k(400 % 3159	
	19
geaaactata otgtgaaca	1.5
+2100+3160	
+210.1 5100 +2115 19	
HILLS DNA	
+:::23 Home sapiens	
The means paptions	
-12L01-	
*.120.* *1123: Cyclin Al ribozyme binding site	
12.0. Cyclin Al Tibozyme binding Site	
-:4005-3160	
	19
acaaqcactt ttggccaga	17
H21CH 3161	
62110-19	
KEILO IDNA	
Color Una	

+:220 + -: -:220 + Cyclin Al ribozyme binding site	
+400 + 3161 baawbacttt tggccagaa	19
+1211 + 3162 +1211 + 19	
HO13+ Bomo sapiens	
+MARR + Cyclin Al ribozyme binding site	
<400 > 3162 aagcactttt ggccagaaa	19
<pre>&lt;:210 &gt; 3163 &lt;:211 &gt; 19 &lt;:212 &gt; DNA &lt;:213 &lt; Homo sapiens</pre>	
<2200 v <2203 Cyclin Al ribozyme binding site	
<pre>&lt;400 + 3163 aqaaacoott gotgoattt</pre>	19
+210.4 3164 +211 = 19 +212 = DNA +213.4 Homo sapiens	
+MAGO+ +MAGO+Cyclin Al ribezyme binding site	
+400> 3164 ttgctgcatt tacagggta	19
+010+ 3165 +011+ 19 +010+ DNA	
<pre>s0130 Homo sapiens colors</pre>	
-MARS Cyclin Al ribozyme binding site	
+:400:2 3165 tgctgcattt acagggtat	19
·(210> 3166	

.

<pre>#211 * 19 #2210 * DNA #2213 * Homo sapiens</pre>	
<pre><u20+ <u23+="" al="" binding="" cyclin="" pre="" ribozyme="" site<=""></u20+></pre>	
H400 + 3166 gotghattta dagggtatt	19
<pre>4210 &gt; 3167 &lt;2211 &gt; 19 &lt;212 &gt; DNA +2213 &gt; Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;323&gt; Cyclin Al ribozyme binding site</pre>	
<400: 3167 ttadagggta ttdattaag	19
+210 + 3168 +211 + 19 +211 > DNA +213 → Homo sapiens	
<pre><din0> <pre><din1.0> <pre>Cyclin A1 ribozyme binding site</pre></din1.0></pre></din0></pre>	
-:400 - 3168 acaqugtatt cattaagtg	19
<pre>&lt;210&gt; 3169 &lt;211&gt; 19 &lt;212&gt; DNA &lt;213&gt; Homo sapiens</pre>	
RODES SOLDS Gyclin Al ribozyme binding site	
-:400:- 3169 caqqqtattc attaagtga	19
+00100+ 3170 +0010+ 19 +0010+ DNA +0013+ Homo sapiens	
+:22:0:- +:22:> Cyclin A1 ribozyme binding site	

0400 · 3170 Agtattoatt aagtgaaat	19
H210 + 3171 H211 + 19 H212 + DMA H213 + H0mb sapiens	
+220 + +223 + Cyclin Al ribozyme binding site	
-(400 + 3171 gtattcatta agtgaaatt	19
6010 + 3172 6011 + 19 6010 + DNA 6013 + Homo sapiens	
<pre><dii0> <di223 -="" al="" binding="" oyclin="" pre="" ribozyme="" site<=""></di223></dii0></pre>	
H400> 3172 aagtgaaatt gtgoottgo	19
<pre>&lt;2108 3173 &lt;211 &gt; 19 &lt;212 + DMA &lt;213 &gt; Homo sapiens</pre>	
<pre>&lt;220 - &lt;223 - Cyclin Al ribozyme binding site</pre>	
k:400 + 3173 attatgoott gootgagtg	19
+1210 s 3174 +1211 s 19 +1212 s DNA +1213 c Homo sapiens	
<pre>&lt;.100 </pre>	
-:400:-3174 gagidagott cataaagog	19
+0010+ 3175 +0010+ 19 +0010+ DNA +0010+ Homo sapiens	

<pre>d200. d200. c200. data</pre>	
High 3175 agragagette ataaagegt	19
+1210 + 3176 +1211 + 19 +1312 + DNA	
-1113 - Homo sapiens	
Hallar Cyclin Al ribozyme binding site	
-:400> 3176 magetteata aagegtaet	19
<pre>&lt;310 &gt; 3177 &lt;3211 &gt; 19 &lt;313 - DNA &lt;313 &gt; Homo sapiens</pre>	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
-:400% 3177 ataaagogta oottgatat	19
+12.10 % 31.78 +12.11 × 19 +12.12 × DNA +12.13 × Homo sapiens	
<pre>schio + schio Supplier Al ribozyme binding site</pre>	
+:400.+ 3178 agogtacott gatatacco	19
+210% 3179 +211% 19 +211% DNA	
+213+ Homo sapiens +226+	
+2230 Cyclin Al ribozyme binding site	
<pre>&lt;:400: 3179 taccttgata taccccatc</pre>	19

+1210 + 3180 +1211 + 19 +1213 + DNA +1213 + Homo sapiens	
<pre>%220 * %223 * Cyclin Al ribozyme binding site</pre>	
+400 + 3180 oottgatata ooccatoga	19
+0010 + 3181 +0011 × 19 +0010 + DNA +0013 + Bimo sapiens	
<pre>&lt;0.00&gt; &lt;0.03&gt; Cyclin A1 ribozyme binding site</pre>	
<400> 3181 ataphopate gacotoage	19
<pre>%210 &gt; 3182 %211 &gt; 19 %212 &gt; DNA %213 &gt; Homo sapiens</pre>	
<pre></pre> <pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
+:400% 3182 categacote agcaagcas	19
<pre>kC10&gt; 3183 +C11&gt; 19 +C1C&gt; DMA +C13&gt; Hemo sapiens</pre>	
+:1200 +:2230 Cyclin A1 ribozyme binding site	
H400H 3183 dcaadcaatt agggagaag	19
+010: 3184 +011: 19 +010: DNA +013: Homo sapiens	
+ 220:223> Cyclin Al ribozyme binding site	

-:400 - 3184 -:aaqmaatta gggagaagt	19
+0210 + 3185 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
<pre>&lt;:220&gt; &lt;:223    Cyclin Al ribozyme binding site</pre>	
-M008 3185 qqqaqaagta caaggotto	19
<pre>&lt;210 * 3186 &lt;2211 * 19 &lt;212 * DNA &lt;213 * Homo sapiens</pre>	
<pre> &lt;220 - &lt;223 - Cyclin A1 ribozyme binding site</pre>	
<400% 3186 tabaaggott baaagtaco	19
<pre>&lt;2108 3187 &lt;2118 19 &lt;2118 DNA &lt;2138 Homo sapiens</pre>	
<pre><d2008 <pre=""><d2038 al="" binding="" cyclin="" pre="" ribozyme="" site<=""></d2038></d2008></pre>	
-:400% 3187 acaaggette aaagtacet	19
+1210 + 3186 +1211 + 19 +1212 + DNA +1213 + Home sapiens	
+mn000 +mn130 Cyclin Al ribozyme binding site	
<pre>+:400% 3188 ottoaaagta ootgtgtgt</pre>	19
+:210:- 3183 +:211:- 19 +:212:- DNA	

+:213 · Homo sapiens	
+1100 +	
+MAB + Cyclin Al ribozyme binding site	
-(400 + 3189)	
tququgtqto cotcatgga	19
H210 + 3190	
8211 × 19	
H212 + DNA	
<pre>&lt;313 * Homo sapiens</pre>	
+1220 +	
+:223 + Cyplin Al ribozyme binding site	
+400 × 3190	
tgtgtccctc atggagcca	19
<210> 3191	
<211 × 19	
SCIENCE DNA	
<pre>%213 / Homo sapiens</pre>	
*(22g)*	
<223> Cyclin Al ribozyme binding site	
<4400 × 3191	19
abotycagtt ottottota	19
<210 * 3192	
<211 + 19	
MOIC + DNA	
+2213 + Homo sapiens	
+:220×	
<pre></pre> <pre></pre> <pre></pre> <pre>C23 &gt; Cyclin Al ribczyme binding site</pre>	
400 - 3192	
delignage to tectedad	19
+110 + 3195	
*III: 19	
-1.11 DNA	
+2130 Homo sapiens	
+2200+ +2230+ Cyclin Al ribozyme binding site	
-22 Crollin Al Tibozyme binding Site	
::4000 3193	
tgcagttctt cttctacaa	19

H210 + 3194 H211 + 19 H212 + DNA H213 + Homb sapiens	
+320 + +3223 + Gyolin Al ribozyme binding site	
HANDE 3194 Geagliteite tictacaat	19
<pre>&lt;210 + 3195 +011 + 19 +0110 + DNA +0013   Homo sapiens</pre>	
<pre>&lt;2205 &lt;2223&gt; Cyclin A1 ribozyme binding site</pre>	
H400+ 3195 amitettett etacaataa	19
<pre>&lt;310   3196 &lt;311   19 &lt;312   DNA &lt;.13   Homo sapiens</pre>	
+CC0+ +CC3+ Cyclin Al ribozyme binding site	
H400% 3196 qttettette tacaataag	19
+1210 + 3197 +1211 + 19 +1212 + DNA +1213 + Homo sapiens	
<pre>summe + summe = Cyclin Al ribozyme binding site</pre>	
0400 - 3197 tottottota caataagtt	19
+0100+3198 +0110+104 +0110+ DNA +00130+ Homo sapiens	
+:22°a>	

-:223 - Cyclin Al ribozyme binding site	
-:4.00 + 31.98	
ttotacaata agtttctga	19
H210 + 31 <b>9</b> 3	
H211 + 19	
1212 + DNA	
<pre><pre></pre></pre>	
<pre>************************************</pre>	
+MARR Syclim Al ribozyme binding site	
+:400 × 3199	
acaataagtt totgaatgg	19
-210 - 220	
+::010 ×   3200 +::011 ×   19	
<pre><pre><c10> DNA</c10></pre></pre>	
<pre>%313 * Homo sapiens</pre>	
1.15 None Supreme	
<210×	
<pre>&lt;223 - Cyclin Al ribezyme binding site</pre>	
<400 ← 3000	
caataagttt ctgaatgga	19
H2105 3001	
(211 - 19)	
*C112: DNA	
<pre>&lt;213&gt; Home sapiens</pre>	
<2208	
-1133 Cyclin Al ribozyme binding site	
<4400N 3201	19
aataagttto tgaatggaa	13
<pre><pre><pre><pre><pre></pre></pre></pre></pre></pre>	
K2118-19	
-CLINDS DNA	
+11130 Home sapiens	
- Trators	
1000 -1000 Cyclin Al ribozyme binding site	
Cyclin hi libozyme binding sice	
-:400:- 3202	
gdaadcactt. ccagaactt	19
+0.10% 3203	
+1.40 + 3203 +22112-19	

+LD12 + DMA +LD13 + Homo sapiens	
*12.20 *	
+CD23+ Cyclin A1 ribozyme binding site	
<pre>-(400 + 3103 quageactic cagaactic</pre>	19
+0103104	
±2211 × 19	
AMO + 2100+	
+1113 + Homo sapiens	
H220x	
<pre></pre> <223 < Cyclin A1 ribozyme binding site	
k:4003204	
todagaadtt daddtodat	19
<010 % 30.05	
<211 × 19	
KÜLÜK DNA	
<pre>&lt;313 - Homo sapiens</pre>	
K2208	
<pre><pre><pre><pre></pre></pre><pre><pre></pre></pre><pre></pre></pre><pre></pre><pre></pre></pre> <pre></pre> <pre< td=""><td></td></pre<>	
+:400:H-3005	
ocadaactic accideata	19
+0.108 3006	
<211> 19	
HOLDS DNA	
+2130 Homo sapiens	
+1m.gs	
+MM38 Cyclin A1 ribozyme binding site	
+:400× 320€	
antinaente catatoaga	19
+m10: 3207	
+D2118-19	
HOLLS DNA	
+213+ Home sapiens	
4:200°	
<pre>%2230 Cyclin A1 ribozyme binding site</pre>	
::400:2-3267	

pappiccata toagaagig	19
H210 + 3208 H211 + 19 H212 + DNA	
20113 - Homo sapiens	
-MLDs - Cyclin Al ribozyme binding site	
<pre>A400 + 3_08 Apthobatato agaagtgoo</pre>	19
HC10+ 3009 HC11+ 19 HC10+ DNA	
G213> Homo sapiens	
<pre>&lt;220&gt; &lt;223&gt; Cyclin Al ribozyme binding site</pre>	
<pre>&lt;:400 &gt; 3209 autgccaata atcgtcata</pre>	19
+02108 3010 +02118 19 +0010 + DMA	
+2213 + Homo sapiens	
+C20+ +C203+ Cyclin Al ribozyme binding site	
-:4008 3210 godaataato gtoataggo	19
+12100 3211 +10110 14	
ROLLO: DMA ROLLO: Homo sapiens	
<pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre>Cyclin Al ribozyme binding site</pre>	
-:400: 3:11 aataatcute ataggette	19
	10
+0.160+3212 +0.110+1+	
+0 125 DNA +02135 Homo sapiens	
talia: nomo saditens	

<220 · 42.23 · Cyclin A1 ribbzyme binding site	
H400 - 3.12 Habbutbata ggottotgo	19
<pre>SUBLE + 3213 SUBLE + 19 SUBLE + DNA SUBLE + Bomb sapiens</pre>	
<pre><pre>&lt;0.000 + &lt;0.000 + Oyolin A1 ribozyme binding site</pre></pre>	
-0400 + 3213 toataggett etgeacgtt	19
<pre>&lt;010&gt; 3214 &lt;011&gt; 19 &lt;010&gt; DNA &lt;013&gt; Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyclin Al ribozyme binding site</pre>	
<4005 3214 cataggette tgcaegttg	19
K210 × 3215 H211 × 19 H212 × EMA H213 × Homo sapiens	
<pre><ildo> <pre><iiii3> Cyclin A1 ribozyme binding site</iiii3></pre></ildo></pre>	
+:4000- 3015 tiptigeabott ggateaact	19
#32100-3316 #32110-13 #32120-INA #32130-Homo sapiens	
+32209 +32200 Cyclin Al ribozyme binding site	
-:400:- 3216 acgttggatc aactaatgt	19
√21n2 3217	

÷211 → 19	
-0.112 · DNA	
<pre><pre><ii113 +="" h="" omo="" pre="" sapiens<=""></ii113></pre></pre>	
4:120 ·	
-CLRB+ Cyplin Al ribozyme binding site	
+(400 + 3.11)	
rdat daadta atgttgttt	19
That had not a egeogeou	
H210 × 3218	
H211 + 19	
<pre><!-- COME</td--><td></td></pre>	
<pre></pre>	
<2207	
<2233 Cyclin Al ribozyme binding site	
<400 > 3218	
aactaatgtt gtttacaat	19
austracyst gereadae	
<210× 3219	
<211 > 19	
<212 > DNA	
<213> Homo sapiens	
•	
K220 K	
<pre>&lt;203 - Cyclin Al ribozyme binding site</pre>	
<400 ← 3219	1.0
taatuttutt tacaatata	19
K010> 3000	
<2115-19	
HOLON DNA	
<pre></pre>	
KD2 () K	
<pre>&lt;0.23 Cyclin Al ribozyme binding site</pre>	
400 0000	
41400 + 3120	19
aatgttgttt acaatatag	19
H210H 3021	
C2119-19	
H2120 ENA	
₹7.00•	
-223> Cyclin Al ribozyme binding site	

-:400 + 3221 - atquigtita daatataga	19
H210 + 3232 H211 + 19 H212 + DNA H213 + Hame sapiens	
<pre>&gt;1220 + &gt;1223 &gt; Cyplin Al ribozyme binding site</pre>	
0400% 3222 gtttacaata tagatgaca	19
+:210 + 3223 +:211 × 19 +:212 > DNA	
<213> Homo sapiens	
<pre>&lt;220&gt; &lt;223 * Cyclin Al ribozyme binding site</pre>	
H400% 3223 ttadaatata gatgadatt	19
<pre>&lt;210 * 3224 &lt;2218 19</pre>	
RCICK DNA RCIBM Homo sapiens	
+220+ +223+ Cyclin Al ribozyme binding site	
<pre>-:400 \ 3004 adaugabatt ttaaaaatg</pre>	19
+0108 3008 +011.+ 19 +0108 DNA	
+Cli35 Home sapiens	
-00200- -00230- Cyclin Al ribczyme binding site	
-:4000- 3.25 qatqacattt taaaaatgt	19
+0100+ 3.26 +0010+ 1+ +0210+ DMA	
+213> Homo samiens	

K220 ·	
-223 - Cyclin Al ribozyme binding site	
+(4)(0 ≠ 3226	
atgadathtt aaaaatgta	19
+:210 + 30:27	
+::::11 + 1.4	
HALL HOMA	
+0.13 · Homo sapiens	
+0.00 ×	
<pre>&lt;.23 * Cyclin Al ribozyme binding site</pre>	
<400 ← 3227	
tgadatttta aaaatgtaa	19
H210> 3228	
<2115 19	
MOION DNA	
RD139 Homo sapiens	
mile deptons	
<0.00×	
<223> Cyclin Al ribozyme binding site	
2400 B000	
<400 × 3228	10
taaaaaatyta aatgaattt	19
<0.10 + 3009	
K211 × 19	
K212: DNA	
H213 Homo sapiens	
+:220x	
-2238 Cyclin Al ribozyme binding site	
+4003 3229	
taaatgaatt tagtttooc	19
au 19 au 10 cu go co co o s	
+2168 3030	
K2115-19	
SOIDE DNA	
+213+ Homo sapiens	
K200)+	
<2230 Cyclin Al ribozyme binding site	
ki400t/ 3230	
aaatgaattt agttteeet	19

HO10 + 3031 HO11 + 19 HO12 + DNA HO13 + Homo sapiens	
N220 - N223 - Cyclin Al ribozyme binding site	
H400+ 3231 satgaattta gttteeett	19
-0010 + 3030 -0011 + 19 -0010 + DNA -0013 - Homo sapiens	
+ AAAO + + AAA> Cyclin Al ribozyme binding site	
<400> 3.3.3 qaatttagtt toocttaga	19
<pre>&lt;210 &gt; 3233 &lt;2210 &gt; 19 &lt;2210 &gt; DNA &lt;2213 &gt; Homo sapiens</pre>	
<220 - <223 Cyclin Al ribozyme binding site	
<pre>&lt;400 + 3133 aatttagttt coottagac</pre>	19
<pre><c108 19="" 3234="" <c110="" <c118="" <c130="" dna="" home="" pre="" sapiens<=""></c108></pre>	
<pre>-dicor</pre>	
<pre>+34600+ 3134 atitagtito cottagact</pre>	19
+0100+ 3:35 +0110+ 19 +0110+ DNA +0130+ Homo sapiens	
+2220:- +223> Cyclin Al ribozyme binding site	

1 1

<pre>%400 + 3235 agtitonith agactitag</pre>	19
-0.10 + 3036 -0.11 + 14 -0.10 + DNA	
-1213 - Homo sapiens	
+:220 + +:223 > Cyclin A1 ribozyme binding site	
<pre>0400 + 3236 gtttocotta gaotttagt</pre>	19
<pre>%210% 3237 %211% 19 %212% DNA</pre>	
<pre>&lt;213&gt; Homo sapiens</pre>	
<pre>&lt;220+ &lt;223    Cyclin A1 ribozyme binding site</pre>	
<400 - 3237 cottagaptt tagtagttt	19
<210 + 3238 <211 + 19 <2210 + DNA	
<pre>AC130 Homo sapiens</pre>	
<pre><d:00> <pre><d:100> <pre>Cyclin A1 ribozyme binding site</pre></d:100></pre></d:00></pre>	
+04008 3238 ottagaettt agtagtttg	19
+00105 3039 +00115 19 +00115 DNA +00135 Home sapiens	
+1200h	
+2233 Cyclin A1 ribozyme binding site	
+:400:- 3239 thagacttta gtagtttgt	19
+:210:- 3::40 +:211:- 19 +:212:- DNA	

Homo sapiens	
+11.20 ·	
H123 - Gyolin Al ribozyme binding site	
-:400 + 3040	
qastttaqta gtttgtaat	19
+:210+ 3241 +:21:+ 19	
HILL TO ANA	
+11.13 · Homo sapiens	
•	
+12.10 +	
-MM3 - Cyclin Al ribozyme binding site	
<400 × 3041	
tttagtagtt tgtaatata	19
K010 × 3042	
KC11 - 19 KC12 - DNA	
<213% Home sapiens	
K220 ×	
<223> Cyclin Al ribozyme binding site	
K400> 3242	
ttagtagttt gtaatatag	19
<pre>&lt;210&gt; 3243 &lt;211&gt; 19</pre>	
<013> DNA	
+00135 Home sapiens	
+:120> +:223> Cyclin Al ribozyme binding site	
allo ijoin ni iidoljme binaing eree	
+(400)+ 3.:43	
ortagtitiqta atatagtoc	19
+0.100+ 30.44	
\$2112-19	
HOLLS DNA	
+11130 Homo sapiens	
\$(2.2.6)	
+22130 Cyclin Al ribozyme binding site	
<pre>44000: 3244 attituata tagtocaac</pre>	19
ulliulaa a lauliida	1 7

+0010 + 3045 +0011 + 19 +0010 + DNA +0013 + Homo sapiens	
-120 - -10.23 - Gyplin Al ribopyme binding site	
+:400.+ 3.145 ttgtaatata gtodaacat	19
<pre>1210 + 3246 1211 &gt; 19 1212 &gt; DNA 1213 &gt; Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyclin Al ribozyme binding site</pre>	
<pre>&lt;400&gt; 3246 taatataqtc caacatttt</pre>	19
+1210 > 3247 +1211 > 19 +1212 > DNA +1213 + Homo sapiens	
<pre>-DDBDBB - Cyclin Al ribozyme binding site</pre>	
<pre>&lt;:400 \cdot 3047 gtdcaacatt ttttaaaca</pre>	19
\$12100 3248 \$12110 19 \$12120 DMA \$1130 Homo sapiens	
<pre><d2000 <="" pre=""> <pre><pre><d2230< pre=""> Cyclin Al ribozyme binding site</d2230<></pre></pre></d2000></pre>	
k4000-3148 todaacatti titaaacaa	19
+0105 3049 +0110 19 +0120 DNA +0130 Homo sapiens	
. 22m.	

·.2200

+CCC3 · Cyclin A1 ribozyme binding site	
+(400 + 3.249 	19
H210 + 3.50 H211 + 1 + H212 + DNA H214 + Homo sapiens	
-MANO - -MANO - Cyclin Al ribozyme binding site	
-:4000-3250 caacattttt taaacaata	19
<pre>%110 x 3251 %011 x 19 %012 x DNA %013 x Homo sapiens</pre>	
<pre>&lt;2208 &lt;223&gt; Cyclin Al ribozyme binding site</pre>	
<pre>&lt;:400&gt; 3251 ascatttttt aaacaataa</pre>	19
+0010 × 3050 +0011 × 19 +0010 × DNA +0013 · Homo sapiens	
<pre><i220.8 <i2235="" al="" binding="" cyclin="" pre="" ribozyme="" site<=""></i220.8></pre>	
-04000 3252 acatittita aacaataaa	19
+0110:- 3253 +0110:- 13 +0110:- ENA +0130: Homo sapiens	
+100s +123> Cyclin Al ribozyme binding site	
-:4003053 -:tawacawta wactgottg	19
<pre>&lt;1110 &gt; 3154 &lt;211 &gt; 19</pre>	

H212 + DNA H213 + Homo sapiens	
+220 + +223 + Cyclin A1 ribozyme binding site	
kung - 3254 tawantgott gtottatga	19
+0210 + 3055 +0211 + 19 +0212 + DMA	
<pre><bl3 <="" <bl30="" homo="" pre="" sapiens=""></bl3></pre>	
<2003 - Cyclin Al ribozyme binding site <4002 3255	
actgottgto ttatgacaa	19
<pre>&lt;210   3256 &lt;211   19 &lt;212   DNA &lt;213   Homo sapiens</pre>	
<pre>&lt;110* &lt;113* Gydlin Al ribozyme binding site</pre>	
<4005 3356 tgcttgtctt atgacaaaa	19
<pre>&lt;210&gt; 3257 &lt;2215 19 &lt;2017   DNA &lt;2115 Homo sapiens</pre>	
-Mids <22038 Cyclin Al ribozyme binding site	
-:4005-3357 gottgtotta tgacaaaaa	19
+0210> 3258 +0011> 19 +0010> DMA +0013> Homo sapiens	
+27.00- +27.80- Cyclin B1 ribozyme binding site	
·:400> 3258	

.

gotgggtgta ggtoottgg	19
<210 + 3.25 3	
×211 + 19	
+1212 + DNA	
+1213 + Homo sapiens	
+1220+	
+:23 + Cyclin B1 ribozyme binding site	
+:400 + 3254	10
quiqtaggto ottggotgg	19
HD10 + 3060	
#211 + 19	
HOLOS DNA	
<pre>&lt;113@ Homo sapiens</pre>	
•	
<0.00%	
<2235 Cyclin B1 ribozyme binding site	
<400> 3260	
gtaggtoott ggotggtog	19
×310× 3321	
<210> 3261 <211> 19	
S212 S2MA	
<213> Homo sapiens	
The state of the s	
K220S	
<223> Cyclin B1 ribczyme binding site	
<4005-3061	
tiqqetggto gggeteegg	19
2015 - 05 CT	
+0108 3262 +0118 19	
HOIDS DNA	
<pre>%2138 Homo sapiens</pre>	
Telis Telis Suprono	
2000S	
-17130 Cyclin B1 ribczyme binding site	
+40m+ 3.16.1	
gategggeten eggtgttet	19
21165 215 C	
+12.10% 316; +12.11% 1 (	
HILLS DMA	
12137 Mama saniens	

+:220 + +:223 + Cyclin B1 ribozyme binding site	
-Mun + 3163 etwoggtytt etgettete	19
+:210 + 3264 +:211 * 19 +:213 + DNA	
Homo sapiens	
+3220 + -3223 + Gyolin B1 ribozyme binding site	
R400% 3264 topggtgtts tgettetes	19
<pre>&lt;210&gt; 3265 &lt;221&gt; 19 &lt;210&gt; DNA</pre>	
+1013> Homo sapiens	
<pre><dable color="block"><dable color="block"><dable color="block"><dable color="block"><dable color="block"><dable color="block"><dable color="block"><dable color="block"><dable color="block"></dable></dable></dable></dable></dable></dable></dable></dable></dable><td></td></pre>	
H400> 3065 tattotgott otocoogot	19
H2108 3266 H2118 19	
+.212> EMA +.213> Homo sapiens	
+:22WS	
+CMM3N Cyclin B1 ribozyme binding site	
+:400:+ 3266 gitatgetta teccegatg	19
+:2108 3067 +:2118 19	
+C21.08 DNA +C21.08 Homo sapiens	
kateas	
+CC2 (> Cyclin B1 ribozyme binding site	
×400× 3267	1 0
totquttota deagetgag	19
2/21 pt. 3/269	

-1211-	19	
-:21.2 -	FNA	
	Homo sapiens	
	•	
<2205		
	Cyclin B1 ribozyme binding site	
-(41)u>	3.168	
	egata egagteace	19
Ca . 49	Symbol Cyaycoacc	± - '
4010s	200	
100 T 100 T		
411.52	Homo sapiens	
2000		
<.11.10 >		
<.1.13>	Cyclin B1 ribozyme binding site	
<400>		
gataaq	gagto accaggaac	19
<210 >		
<211>		
<212>		
<213>	Homo sapiens	
<220%		
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<400>		
ocagga	lacto gaaaattaa	19
<210>		
<2115		
4212×		
<213 %	Homo sapiens	
SJ 2008		
<2223.5	Cyclin B1 ribozyme binding site	
- [4 ()(1)	3271	
ot.ogaa	laatt aatgotgaa	19
-12105-		
::211:\		
.:212:-	ENA	
·:21:::-	Homo sapiens	
-:220C:-		
·12231-	Cyclin Bl ribozyme binding site	

<pre>(44)) + 3272 toguaaatta atgotgaaa</pre>	19
H210 + 3273 H211 + 19 H212 + DNA H213 + Hamb sapiens	
+0000 + +0003 + Cyclin B1 ribozyme binding site	
-:400 × 3273 gotyaaaata aggogaaga	19
<pre>&lt;210 + 3274 +211 &gt; 19 &lt;212 + DNA &lt;213 + Homo sapiens</pre>	
<pre><dii0 -="" <ili=""><li><pre></pre> <pre></pre> <pre></pre></li></dii0></pre>	
त्400 × 3274 व्यव्यवस्थानम् aacatggca	19
<pre>KD10 = 3275 HD11 + 14 HD12 + DNA HD13 + Homo sapiens</pre>	
<pre>&lt;220 - &lt;223   Cyclin B1 ribozyme binding site</pre>	
+:400 + 3275 aaaqogoqtt cotaoggoo	19
+12108 3.76 +1211 - 19 +1212 DNA +1213 Homo sapiens	
-1220- -12138 Cyclin Bl ribozyme binding site	
+400+ 3176 auguqugtto ctaeggees	19
+02100+ 3077 +0210+ 14 +0210+ DNA +0210+ Homo sapiens	

H223 · Cyclin B1 ribozyme binding site	
H400% 3277  opequationta oggenootg	19
+0010 + 3078 +0011 + 19	
HUIL - DNA HUIL - Homo sapiens	
<pre>&lt;:220&gt; &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
<400> 3278 otgoaacoto caagooogg	19
<210> 3279 <211> 19 <212> DNA	
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<pre>&lt;210&gt; 3280 &lt;211&gt; 19 &lt;212&gt; DNA</pre>	
<213> Homo sapiens	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
k4005 3180 aacagotott ggggacatt	19
+00100+ 3081 800108-19	
80108 DMA 80130 Hemo sapiens	
<pre>&lt;22000 &lt;22030 Cyclin B1 ribozyme binding site</pre>	
-:400:-3::81 tggggacatt ggtaacaaa	19

H2105 (			
·213 · I			
	iomo sapiens		
KI220 +			
	Cyclin Bl ribozyme	binding	site
		Ĭ	
-(4)0(	31.81		
gabatt	ggta acaaagtca		1:
-121n - 1			
<pre>&lt;0111 &gt; 1</pre>			
S213 - I	Homo sapiens		
<330°			
	Cyclin B1 ribozyme	hinding	cito
Server Description	.yciin bi fibozyme	binaing	Site
<4005 3	9783		
	agto agtgaacaa		1:
	.,,.,		
ki210 × 3	3284		
<211 > 1			
42125 E	PNA		
<2135 E	Homo sapiens		
H220%			
K2235 (	Cyclin B1 rib:zyme	binding	site
	2001		
<400 × 3			19
aaaayi	octa tgaagaagg		1.
40 10 × 3	\$ 18 K		
-111-1			
kir 12 × 0			
	Homo sapiens		
	•		
-12201-			
-12231- 0	Cyclin B1 ribozyme	binding	site
- (4000 - 3			
qcaaaaa	eett cagetactg		19
,	t v o v		
-01000 3			
::711:- 1 ::712:- 5			
	ла Lomo sapiens		
	Namo pahiena		
-12201-			
	Cyclin B1 ribozyme	bindina	site

-:400 + 3186 basascotto agotactgg	19
+0010 + 3087 +0011 + 19 +0012 + DNA +0013 + Homo sapiens	
+323 Gyolin Bl ribozyme binding site	
-400 × 3287 wottbagata otggaaaag	19
<pre><c10> 3288 <c11> 19 <c12> DNA <c13> Homo sapiens</c13></c12></c11></c10></pre>	
<pre><dmodel <pre=""><dmodel< pre=""> <pre><dmodel< pre=""> <pre><pre><pre><pre></pre> <pre></pre> <pre><pre><pre></pre> <pre></pre> <p< td=""><td></td></p<></pre></pre></pre></pre></pre></dmodel<></pre></dmodel<></dmodel></pre>	
<pre><d400% 3088="" attgataaa<="" pre="" tqqaaaaqtc=""></d400%></pre>	19
+:010 + 3089 +:011 > 19 +:010 > DNA +:013 > Homo sapiens	
<pre>&lt;2208 &lt;2239 Gyolin B1 ribozyme binding site</pre>	
-:400% 3289 aasagtoatt gataaaaa	19
+12108 3190 +12118 13 +1212: DNA +1213: Homo sapiens	
-2200- -2230 Cyclin B1 ribozyme binding site	
-:400:-3190 gtoattgata aaaaactac	19
00100-3091 00100-19	

<pre><!--213 • Homo sapiens</pre--></pre>	
<pre><did0 <pre=""><did3cyclin b1="" binding="" pre="" ribozyme="" site<=""></did3cyclin></did0></pre>	
<pre>-:400 + 3291 tagagaacta ccaaaacct</pre>	19
<pre>#AB10 + 3A92 +AB11 + 19 #AB12 + DNA +AB13 + Homo sapiens</pre>	
RC20 - RC23 - Cyclin B1 riboxyme binding site	
<400 · 3192 ocaaaaosto ttgaaaagg	19
<pre>&lt;210% 3293 &lt;211% 19 &lt;212% DNA &lt;213% Homo sapiens</pre>	
<220> <223> Cyclin B1 ribozyme binding site	
<400% 3193 aaaacotott gaaaaggta	19
<pre>\$210&gt; 3294 \$211&gt; 19 \$212&gt; DNA \$22130 Homo sapiens</pre>	
<pre>scides scides Gydlin B1 ribbzyme binding site</pre>	
k4005-3294 tqaaaaggta ootatgotg	19
+02100+ 3295 +0211> 19 +02100+ DMA +02130+ Hemo sapiens	
<pre>+C2008 +C2230 Cyclin B1 ribozyme binding site</pre>	
+:400:- 3195	19

H210 + 3296 H211 + 19 H213 + DNA H213 + H2mo sapiens	
<pre>+M20+ +M23+ Gyolin B1 ribozyme binding site</pre>	
<pre><i400% 3096="" pre="" tgagccagt<="" twocagtqto=""></i400%></pre>	19
+1.10 + 3197 +1211 + 19 +1210 > DNA +1212 > Homo sapiens	
<pre>shhos shhos Cyclin B1 ribozyme binding site</pre>	
(400) 3097 tgagcotgtt aaagaagaa	19
+2105 3198 +2112 19 +2125 DNA +213 - Homo sapiens	
+:220 * +:223 * Cyclin B1 ribbzyme binding site	
k4008 3298 gaqootgtta aagaagaaa	19
+0100+ 3199 +0210+ 19 +0210+ DNA +0213> Home sapiens	
-ON OF- -ONESC Cyclin Bl ribozyme binding site	
H400> 3194 agaaaaacti togootgag	19
+2210+ 3300 +2210+ 19 +2210+ DNA +22130+ Homo sapiens	

4220>

<pre><!--!!!! Cyclin B1 ribozyme binding site</pre--></pre>	
H400+ 3300 gaaaaantt ogootgago	19
<pre>%210 + 3301</pre>	
kCLO+ kDD3+ Cyclin Bl ribozyme binding site	
<pre>&lt;400 + 3301 aaaaacttto gootgagee</pre>	19
<pre>&lt;0.10 &gt; 3302 &lt;0.11 &gt; 19 &lt;0.10 &gt; DNA &lt;0.13 &gt; Homo sapiens</pre>	
<220> <220> <220>	
<400> 3302 cotqagocta ttttggttg	19
<pre>kD10 &gt; 3303 </pre> <pre><d11> DNA </d11></pre> <pre>kD13 &gt; Home sapiens</pre>	
k2208 k2239 Gyolin B1 ribozyme binding site	
k4008-3303 tuaqeetatt ttggttgat	19
+1110+ 3704 +1111> 19 +1110+ DNA +1113> Homo sapiens	
-(2000) -(2030) Cyclin Bl ribozyme binding site	
::4000- 3:04 qagootaftt tggttgata	19
-1.:10+-33:05 -:2112-19	

4010 - DNA 4013 - Homo sapiens	
<pre>0000 + 00003 + Cyclin B1 ribozyme binding site</pre>	
H400 - 3305 apporatitt ggttgatac	19
80210 + 3306 +0211 + 19 +0212 + DNA +0213 - Homo sapiens	
<pre>&lt;220&gt; &lt;123&gt; Cyclin B1 ribozyme binding site</pre>	
<400> 3306 tatittggtt gatactgcc	19
<pre>&lt;210&gt; 3307 &lt;211&gt; 19 &lt;110&gt; ONA &lt;113&gt; Homo sapiens</pre>	
K220> K2238 Cyclin B1 ribozyme binding site	
<pre>-:400&gt; 3307 ttggttgata_ctgcctctc</pre>	19
<pre>&lt;0.105 3308 +00.105 19 +00.105 CMA &lt;00.130 Homo sapiens</pre>	
<pre>&lt;:220:- &lt;:223:- Cyclin B1 ribozyme binding site</pre>	
+:4005 3308 atactgooto tocaagooo	19
KCC108 3309 +CC110+ 19 +CC1C08 ENA +CC1309 Home sapiens	
+22100+ +22130+ Cyclin B1 ribczyme binding site	
+:400% 3:09	

astgoctsts caageccaa	19
+1210 + 3310 +1211 + 13 +1212 + DNA +1213 + Homb sapiens	
<pre>+320 - +323 - Cyclin B1 ribozyme binding site</pre>	
ਲ400 + 9310 tqqaaacato tggatgtgo	19
<pre>KC10 × 3311 KC11 × 19 KC12 &gt; DNA KC13 × Homo sapiens</pre>	
<1.00% <1.03% Cyclin B1 ribozyme binding site	
<pre>&lt;400x 3311 qacctgtgtc aggctttct</pre>	19
<pre>#310% 3312 #211% 19 #211% ENA #213% Homo sapiens</pre>	
+(220) +(223) Cyclin B1 ribozyme binding site	
<pre>k:400 \ 3312 tgtcaggott tototgatg</pre>	19
+210 + 3313 +211 \ 19 +212 \ 2MA +2013 \ Home sapiens	
-thus -thus Cyclin B1 ribozyme binding site	
<pre>44UU-3313 gtcadgettt otetgatgt</pre>	19
<pre>&lt;:210 + 3:14 &lt;:211 + 1* &lt;:211 + DNA &lt;:215 &gt; Homo sapiens</pre>	

<pre>+:2.00 + +:2.03 + Oyclin B1 ribozyme bind</pre>	ding site	
H400 - 3314 toaggottto totgatgta		19
+0010 + 3315 +0011 + 19 +0010 + DNA +0015 + Homo sapiens		
<pre><d200 -="" <d203="" bind<="" bl="" gyolin="" pre="" ribbzyme=""></d200></pre>	ding site	
<pre>&lt;400% 3315 aggotttoto tgatgtaat</pre>		19
+:210 + 3316 4:211 > 19 +:212 > DNA +:213 > Homo sapiens		
<pre>&lt;:220 * ::223 * Cyclin B1 ribbzyme bind</pre>	ding site	
<pre>-:4008 3316</pre>		19
+3210.5 3317 +3211.5 19 +32125 DNA +32135 Homo sapiens		
-0.000- -0.000- Gyolin El ribozyme bind	ding site	
-:400:- 3317 tgatgtaatt ottgoagta		19
<pre>%0100 3318 %0110 19 %0110 DNA %0180 Homo sapiens</pre>		
<pre>*MRIGH* *MRIFF* Cyclin El ribozyme bind</pre>	ding site	
-(400)- 3518 gatqtaatto ttgcagtaa		19
+ 2165 3319		

-::111 ·	19		
·::12 ·			
	Homo sapiens		
1	Timo Supremo		
-1227-			
	Cyclin Bl ribozyme	hindina	sita
	o, or in Bi iibozyme	Dinaing	3100
·(40) ·	2.10		
			1.0
Lutada	ttott gcagtaaat		19
2000	5.5.5.5		
1.10			
<111			
<21.15			
4213 ·	Homo sapiens		
<220 ×			
<2233	Cyclin B1 ribozyme	binding	site
<400>			
tattq	caqta aatgatgtg		19
<210 >			
<11>	19		
<2112 >	AUC		
<21.35	Homo sapiens		
400008			
<2223 ×	Cyclin B1 ribozyme 1	binding	site
-14000	3321		
ggaget	igato caaacettt		19
	-		
-121,68	3302		
2011S			
	Homo sapiens		
	THE CAPTOR		
02268			
	Cyclin B1 ribozyme 1	hindina	site
	tysiin bi iibsiyme .	~ 1	
-:4005	3300		
	accit tgtagtgaa		19
.vaac	ast. c egeagegaa		1 2
-121 01-	च ४७ <del>व</del>		
Translation (Co.)			
ts Elizabet	Homo sapiens		
and the second			
*(21,0)*	a .11. p1 11	1- 11 1	-1
. //	Cyclin Bl ribozyme N	binaina -	SITE

<pre>&lt;400+ 3323 &gt;caaacottt gtagtgaat</pre>	19
+0210 + 3324 +0211 + 19 +0012 + DMA +0013 + Homo sapiens	
+C220 + +C223 + Cyclin B1 ribozyme binding site	
<pre>k400 + 3304 aacotttqta gtgaatatg</pre>	19
<pre>&lt;010 &gt; 3305 &lt;011 &gt; 19 &lt;012 &gt; DNA</pre>	
<pre>&lt;313/ Homo sapiens</pre>	
<pre>&lt;320 * &lt;123 * Cyclin B1 ribozyme binding site</pre>	
<400 - 3325 gtagtgaata tgtgaaaga	19
<210 > 3306 <211 > 19 <212 > DNA	
<213  Homo sapiens	
<2200: <223> Cyclin B1 ribozyme binding site	
x400% 3306	19
qtqaaagata tttatgctt	13
+:210% 3327 +:211% 19	
HIGHER ENA	
-1111 to Homo sapiens	
<pre>STATES Cyclin B1 ribozyme binding site</pre>	
H400H 3327	
daaaqatatt tatgottat	19
+:210:+ 3328	
+0210-14	
<pre>+:21:&gt; DNA +:21:&gt; Home sapiens</pre>	
ero. Himo pupitono	

<pre>000000 000000000000000000000000000000</pre>	
<pre>&lt;400 + 3328 aaaqatattt atgettate</pre>	19
<pre><di10 +="" 3329="" <pre=""><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></di10></pre>	
RC13 - Homo sapiens	
<pre><pre><pre><ill0 +="" -="" <ill23="" b1="" binding="" cyclin="" pre="" ribozyme="" site<=""></ill0></pre></pre></pre>	
<400 · 3309 aagatattia tgottatot	19
<pre>&lt;010 + 3530 &lt;011 &gt; 19 &lt;012 &gt; DNA &lt;013 &gt; Homo sapiens</pre>	
<pre>&lt;320 - &lt;323 - Oydlin B1 ribbzyme binding site</pre>	
<pre>+(400 + 3330 atttatgett atetgagae</pre>	19
<pre><pre>&lt;010 + 3331 &lt;011 &gt; 19 &lt;012 &gt; DNA</pre></pre>	
<2135 Homo sapiens	
<pre><pre><pre><pre><pre></pre> <pre><pre></pre> <pre><pre></pre> <pre></pre> <pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre></pre> <p< td=""><td></td></p<></pre></pre></pre></pre></pre></pre></pre>	
<pre>&lt;4000-3331 titatgetta totgagada</pre>	19
+1110+ 3332	
H011H 19 H012H DNA	
+20133 Homo sapiens	
+:210::213:- Cyclin Bl ribozyme binding site	
<pre>&lt;:400: 3332 tatqcttatc tgagacaac</pre>	19

0010 + 3333 0011 + 14 +010 > DNA +013 + Homo sapiens	
<pre><ul><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton</li><li>culton<!--</td--><td></td></li></ul></pre>	
<pre>6400 + 3333 gagaaaatt gaggaagag</pre>	19
+0210 + 3334 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
<pre>&lt;:220 &gt; &lt;:223 &gt; Cyclin B1 ribozyme binding site</pre>	
<pre>%400% 3334 qcaaqcaqtc agaccaaaa</pre>	19
<pre>&lt;210&gt; 3335 &lt;211&gt; 19 &lt;212&gt; DNA &lt;213&gt; Homo sapiens</pre>	
+3205 +323 - Cyclin B1 ribozyme binding site	
-:400:- 3335 qaddaaaata dotactggg	19
+1210+ 3336 +1211+ 19 +1212+ 5NA +1213+ Homo sapiens	
+0.2008 +0.030+ Cyclin El ribozyme binding site	
-:400:- 3336 Aaaataccta ctgggtcgg	19
+1210: 3337 +1211: 19 +1211: DNA +1213: Homo sapiens	
<pre>+0.00&gt; &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	

÷(40) ÷ 3337	
otwotggate gggaagtea	19
(21.0	
+:210 × 3338	
-12115 19	
HILLS DNA	
-1213 - Homo sapiens	
<208	
Hullas Cyclin B1 ribozyme binding site	
The system of the state of the	
+:400 / 3 / 38	
togggaagto actggaaac	19
<pre><pre><pre></pre></pre></pre>	
±111> 19	
HO12> DNA	
+2213> Homo sapiens	
+12.20×	
+10.03> Cyclin B1 ribozyme binding site	
+:400> 3339	
gagagocato otaattgac	19
gagagecase ecaacegae	+ -/
HILLUS 3340	
<1115-19	
HOLLES DNA	
+2213> Home sapiens	
·	
+2233 Gyolin El ribozyme binding site	
+(4)(i) x - 3;(40)	
aqodatoota attgactgg	19
H0109 3341	
40015-19	
HONDONA	
+11130 Homo sapiens	
- Carriers	
+CLLS+ Cyclin B1 ribozyme binding site	
+:4000+ 3341	
datoctaatt gactggcta	19
- Auto-Sociation gaologyoca	1.7
H210H 3342	
±211:- 19	
-21'n, FeID	

<pre><!--213 + Homo sapiens</pre--></pre>	
<pre>&lt;:220 • &lt;:220 • Cyclin B1 ribozyme binding site</pre>	
H400 + 3:42 tgantggota gtacaggtt	19
+.210 + 3743 +.211 + 19 +.212 + DMA +.213 + Homo sapiens	
+330 + +333 + Cyclin B1 ribozyme binding site	
+400 × 3343 ctapctagta caggiticaa	19
<pre>*210 * 3344 *211 * 19 *212 * DNA *213 * Homo sapiens</pre>	
+220% +22%+ Cyclin B1 ribozyme binding site	
+400 + 3344 aqtacaggtt caaatgaaa	19
<pre>&lt;.010 &gt; 3345</pre>	
-220 - -223 - Cyclin B1 ribozyme binding site	
+400+ 3345 gtadaggtto aaatgaaat	19
+010+ 3346 +0115 19 +010+ DNA +013+ Hemo sapiens	
+22co Cyclin Bl ribozyme binding site	
+40C: 3346 aaatgaaatt caggttgtt	19

<pre>&lt;:210 &gt; 3347 :211 &gt; 19 :2112 &gt; DNA :2113 &gt; Homo sapiens</pre>	
<pre><pre><pre><il20 *="" <li=""><pre>Cyclin B1 ribozyme binding site</pre></il20></pre></pre></pre>	
+:400 × 3347 aatgaaatto aggttgttg	19
HC10 + 3548 HC11 + 19 HC10 + DMA HC13 * Homo sapiens	
<pre>&lt;220 &gt; &lt;223 · Cyclin B1 ribozyme binding site</pre>	
-:4003348 Battcaggtt gttgcagga	19
+0210 × 3349 +0211 × 19 +0212 > DNA +0213 + Homo sapiens	
<pre>2000 - 3003 Cyclin B1 ribozyme binding site</pre>	
<400> 3349 toaggttgtt gcaggagac	19
<pre>%2108 3350 %2118 19 %2128 DNA %2138 Homo sapiens</pre>	
+MLEOS +MLEOS Cyclin B1 ribozyme binding site	
<pre>&lt;4000 3350 agaccatqta catgactgt</pre>	19
+0.10% 3351 +0.11% 19 +0.11% DNA +0.1% Homo sapiens	
·(229)	

-:233+	Cyclin Bl	ribozyme	binding	site		
0400 + Datqa¢	3351 tato toda	ittatt				19
+0010 + +0011 + +0017 +	1.9					
	H⊙m∋ sapi	ens				
-:220 · -:223 ·	Oyalin Bl	ribozyme	binding	site		
<400 / tgabtg	3352 toto datt	attga				19
<210×	19					
	UGA Homo sapi	ens				
<:220> <:223>	Cyclin B1	ribozyme	binding	site		
<400> tgtata	3353 datt attg	atcgg				19
<210 s	19					
<212 × < < 113 × < < < 1 × < < < < < < < < < < < < < <	DNA Homo sapi	ens				
<0000.+ <0003.+	Oyolin Bl	ribozyme	binding	site		
+(400)≥ gtotoc	3354 atta ttga	teggt				19
+:::10:+ +::11:- +::11::-	19	0.7.0				
::22(0):	пошь зарі	E112				
	Cyclin B1	ribozyme	binding	site		
-14000 otlogat	NA5: tatt gato	ggttc				19
·1710:-						

HOME Sapiens	
+:223 + Cyclin B1 ribozyme binding site	
<pre>0400 + 3356 attattgato ggttoatgo</pre>	19
H210 - 3357 H211 - 19 H211 - DNA H213 - Homo sapiens	
<pre>&lt;22005 &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
<400> 3357 ttgatoggtt catgoagaa	19
<pre>&lt;210* 3358 &lt;2212* 19 &lt;2212* DNA &lt;2213* Homo sapiens</pre>	
+1100 +1113 × Cyclin B1 ribozyme binding site	
<pre>04008 3358 tgateggtte atgeagaat</pre>	19
+210% 3359 +211% 19 +212% DNA +213% Homo sapiens	
<pre>+mm0&gt; +mm0&gt; +mm0 Strip Cyclin El ribozyme binding site</pre>	
H400H 3359 atgcagaata attgtgtgc	19
+010: 3360 +011: 13 +010: DNA +013: Homo sapiens	
+:2::0:- +:2::3:- Cyclin B1 ribozyme binding site	
+:400× 3560	

cagaataatt gtgtgooca	19
+17.10 × 33.61	
±1211 × 13	
H212 DNA	
H2130 Homo sapiens	
•	
+(220)*	
<pre>+1223 ** Cyplin B1 ribozyme binding site</pre>	
+:400 + 3361	
quaquitqqtt ggtgtcact	19
131A. 35C	
+:210 × 3367 +:211 × 19	
+21.0 DNA	
H2139 Homo sapiens	
- Lib Hame Out Land	
ki220>	
<pre>&lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
+400> 3362	
ggttggtgtc actgccatg	19
<210 × 3363	
<211> 19	
RILLIAN ENA	
RC2135 Homo sapiens	
+(210 s	
+2213> Cyclin B1 ribozyme binding site	
The state of the s	
+:4008-3363	
ctgccatgtt tattgcaag	19
:0.100 3364	
H111: 19	
HOLLS DNA	
+2130+ Homo sapiens	
+:220:-	
+223% Cyclin El ribozyme binding site	
Selection of the Districting Side	
+(400): 3364	
tgcdatgttt attgcaage	19
+CD100+ 3:465	
<2.11 (-1.)	
K21.DE DNA	
<213> Homo sapiens	

4220 -			
<pre>&lt;:223 · Cyclin B1 ribozyme</pre>	binding	site	
-(4.5.) - 33.65			
gobatgitta tigoaagoa			19
+210 + 3366			
4211 + 19			
HITT - DNA			
<pre>&lt;0.013 - Homo sapiens</pre>			
CONTRACTOR DESCRIPTION	1 1 11	• ,	
+2223 - Cyclin B1 ribozyme	binding	site	
<400 + 3366			
catqtttatt gcaagcaaa			19
<010 × 3367			
<211 / 19			
<312 > DNA			
k213 - Homo sapiens			
<220 ·			
<2223 - Cyclin B1 ribozyme	binding	site	
	_		
<400 × 3367			
daagdaaata tgaagaaat		-	19
<210 ± 3368			
<211> 19			
<0105 DNA			
<213> Homo sapiens			
<.220 s	1 1 1 1		
<pre>SDD35 Cyclin B1 ribozyme</pre>	binding	Site	
<400 × 3368			
aagaaatgta cootocaga		<u>-</u>	19
KL105 3369			
-00116 19			
<pre><cli>Clic DNA</cli></pre> <pre><cli>Clic Homo sapiens</cli></pre>			
all we now sapiens			
kt <u>2</u> 17 (d) k			
-2233 Cyclin B1 ribozyme	binding	site	
<:400:- 3369 			
atgracecte cagaaattg		1	19
·.210:- 3370			

<pre>4011 + 13 4012 + DNA 4013 + Homo sapiens</pre>	
-:220:223 - Cyclin B1 ribozyme binding site	
+(400 + 3070) topagaaatr ggtgacttt	19
+1210 + 3371 +1211 + 10 +1211 + DMA +1213 + Homo sapiens	
K220) K223% Cyclin B1 ribozyme binding site	
+:4005 3571 ttggtgact: tgcttttgt	19
+1210 3372 +1211 × 19 +1212 × DNA +1213 · Homo sapiens	
+0000 +0000 Cyclin B1 ribozyme binding site	
<pre>&lt;400&gt; 3372 tggtgacttt gottttgtg</pre>	19
+3210> 3373 +3211> 14 +3312> DNA +3313> Homo sapiens	
+3208 +3213> Cyclin B1 ribozyme binding site	
<pre>c4f0x 3373 quotitigett tigtgactg</pre>	19
+D108 3374 +D118 14 +CD120 DNA +D130 Homo sapiens	
<pre><pre><t220> <t223> Cyclin B1 ribozyme binding site</t223></t220></pre></pre>	

-:400 + 3374 Abhittgoitt   tgtgactga	19
+1210 + 3375 +1211 + 19 +1212 + DNA +1213 + Homo sapiens	
+0.00 + +0.00 + Oyolin B1 ribozyme binding site	
<pre>&lt;:400 + 3376 atttqctttt gtgactgac</pre>	19
+210% 3376 +211% 19 +213% DNA +213% Homo sapiens	
+0.008 +0.008 +0.003 Cyclin Bl ribozyme binding site	
<pre>44008 3376 gacaacastt atactaage</pre>	19
<pre><pre><pre><pre></pre></pre> <pre><pre></pre> <pre></pre> <pre< td=""><td></td></pre<></pre></pre></pre>	
<pre><pre><pre><pre></pre></pre></pre></pre>	
<pre>&lt;:223&gt; Cyclin B1 ribozyme binding site &lt;:4005 3377</pre>	
adaadadta tadtaagda	19
<pre>#210# 3378 #2118 19 #2128 2MA #2128 Homo sapiens</pre>	
<pre>%TIOP %TION Cyclin B1 ribozyme binding site</pre>	
+400% 3378 aacacttata ctaagcacc	19
+11 1(:- 3379 +1111: 19 +1111: DNA +1213> Homo sapiens	

<pre>+:220 + +:223 * Cyclin B1 ribozyme binding site</pre>	
+(400 + 3379 antitatauta agcaccaaa	19
+2210 + 3580 +2211 + 19 +2212 + DNA +2213 + Homo sapiens	
<pre><i2.0 +="" -="" <i2.3="" binding="" bl="" gyolin="" pre="" ribozyme="" site<=""></i2.0></pre>	
2400 - 3380 gbabbaaatb agacagatg	19
+210 + 3381 <211 + 19 +212 ENA <213 Homo sapiens	
<pre>+220&gt; &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
<pre>+:400 + 3381 aatgaagatt ctaagaget</pre>	19
+010+3382 +211+19 +212> DNA +213> Homo sapiens	
-:220 -:223 - Cyclin Bl ribozyme binding site	
-400-3382 atgaagatto taagagott	19
<pre>%2100+ 3383 %2110+ 19 *22120+ DNA *2213+ Homo sapiens</pre>	
(2.70) (2.70) Cyclin Bl ribozyme binding site	
H41UP 3383 gaagattota agagottta	19

+:210 + 3:884 +:211 + 14 +:212 + DNA +:213 + Homo sapiens	
-:120: -:123 - Cyclin B1 ribozyme binding site	
<pre>+3400 + 3384</pre>	19
+210% 3385 +211 + 19 +212% DNA +213% Homo sapiens	
<pre>&lt;:220 * &lt;:223 * Cyclin B1 ribozyme binding site</pre>	
+:400 > 3385 taaqagettt aaaetttgg	19
\$210 + 3386 \$211 + 19 \$212 + DNA \$213 * Homo sapiens	
<pre>+CMCO &gt; +CMCO &gt; +</pre>	
<pre>x400 + 3586 aagaqcttta aactttggt</pre>	19
+210 + 3387 +211 > 19 +212 > DNA +213 > Homo sapiens	
+:220+ +:223+ Cyclin B1 ribozyme binding site	
-:4003387 otttaaactt tggtotggg	19
+1110+ 3388 +111+ 19 +1110+ DNA +1113+ Homo sapiens	
<pre>4.200 </pre> <pre>Cyclin B1 ribozyme binding site</pre>	

-:400 + 3388 tttaaacttt ggtctgggt	19
+1210 + 3389 +1211 + 19 +1212 + DNA +1213 + Homo sapiens	
+:220.4 +:223.4 Cyclin B1 ribozyme binding site	
+:400 + 3389	
aastitggts tgggtegge	19
<pre>&lt;210&gt; 3390 &lt;211&gt; 19 &lt;212&gt; DMA &lt;213&gt; Homo sapiens</pre>	
<220 · <223 · Cyclin B1 ribozyme binding site	
:400 × 3390 gqtctggqtc ggcctctac	19
<pre>%210 + 3391 %211 + 19 %212 + DNA %213 * Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyplin Bl ribozyme binding site</pre>	
44009 3391 agtaggeate tacettige	19
+210 × 3332 +211 + 19 +211 > DNA +213 > Econo sapiens	
₹#208	
+:0230 Cyclin B1 ribozyme binding site	
4400% 3391 togacctota cottigoac	19
+02100+ 33335 +0211+ 19 +0212+ DNA	

<pre>+1213 * Homo sapiens</pre>	
+:220 +	
+:223 - Cyclin B1 ribbzyme binding site	
-:400 + 3393	
detendent typeactics	19
+:210 +   3394	
+1211 + 19	
+121.0 + DMA	
+1213 + Homo sapiens	
×1220 ×	
-:223 - Cyclin B1 ribozyme binding site	
+400+ 3394	
ototacetti geaetteet	19
<210 + 3395	
<211 > 19	
K212× DNA	
+213 - Homo sapiens	
+0.00 ×	
<pre>SCACS Cyclin B1 ribozyme binding site</pre>	
K400 + 3395	
etitqcaett cottoggag	19
K210 × 3396	
<2115-19	
HIDIDA DINA	
+215 Homo sapiens	
<(20)÷	
+:173 - Cyclin B1 ribozyme binding site	
+:400:- 3296	
ttignasite ottoggaga	19
H210H 3397	
+IIII: 19	
+01120 DNA	
+Clis-Homo sapiens	
Righter	
RELATE Cyclin Bl ribozyme binding site	
÷4000 3397	
gcacttcctt cggagagca	19

+0010 + 3398 +0011 + 19 +0010 + DNA +0013 + Homo sapiens	
<pre><pre><pre>&lt;1.20 * &lt;1.23 * Gyolin B1 ribozyme binding site</pre></pre></pre>	
K400 + 3398 capticotto ggagagoat	19
<pre>SU10 + 3399 SU11 + 19 SU13 + DMA SU13 + Homo sapiens</pre>	
<pre>&lt;000&gt; &lt;0008 Cyclin B1 ribozyme binding site</pre>	
k400 = 3399 ggagagcatc taagattgg	19
<pre>&lt;210% 3400 &lt;211% 19 &lt;213% DNA &lt;213% Homo sapiens</pre>	
+2235 Cyclin B1 ribozyme binding site	
र400> 3400 adadcatota agattggag	19
+2108 3401 <2118 19 <2118 DNA <2130 Homo sapiens	
<pre>*Vilus *Vilus* Cyclin B1 ribozyme binding site</pre>	
k400% 3401 arctwagatt ggagaggtt	19
+12100+ 3402 +12110+ 19 +12110+ DNA +12120+ Homo sapiens	
+:22a>	

-:223 · Cyclin B1 ribozyme binding site	
H400 - 3402 tggagaggtt gatgtcgag	19
+1210 + 3403 +1211 + 19 +1212 + DNA +1213 + Homo sapiens	
+0000+ +0003+ Cyclin B1 ribozyme binding site	
k400 + 3403 qqttqatqto gagcaacat	19
<pre>&lt;010 × 3404 &lt;011 &gt; 19 &lt;012 &gt; DNA &lt;013 &gt; Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
<400> 3404 qaqcaacata ctttggcca	19
<pre>&lt;0.10 &gt; 3405 &lt;0.11 &gt; 19 &lt;0.17 &gt; DNA &lt;0.13 &gt; Homo sapiens</pre>	
<pre>%2208 %213% Syclin B1 ribozyme binding site</pre>	
0400% 3405 daadatadtt tggddaaat	19
+:210+ 3406 +:211> 19 +:217> DMA +:213> Homo sapiens	
-17200 -1723- Cyclin B1 ribozyme binding site	
-:400:- 3406 aacatactit ggccaaata	19
+210:- 3407 +211:- 19	

+1312 + DNA	
+13 · Homo sapiens	
i ) I mmo saptens	
3.1.2.0 -	
-1.123 * Cyclin B1 ribozyme binding site	
+:400 / 3407	
tuqobaaata ootgatgga	19
-:210 + 3408	
×2211 × 19	
HIZIL > DNA	
+113 - Homo sapiens	
K120 A	
<pre>&lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
·400 · 3408	
gatggaacta actatgttg	19
+:210 + 3409	
K211 + 19	
<pre>&lt;212 &lt; DNA</pre>	
-213 · Homo sariens	
<:220 ×	
<223 - Cyclin B1 ribozyme binding site	
Selv Systim Di Tibolymo binaing bios	
+:40p + 3409	
gaadtaadta tgttggadt	19
and sudamed to get eggete	
<210 = 3410	
*2.118 19	
K212S DNA	
s213% Homo sapiens	
N.2132 Hould Saptems	
K11.008	
<pre>&lt;0.03% Cyclin B1 ribozyme binding site</pre>	
- 1500 - 0.110	
\$14.00(s) 34.10	1.0
taketatett ggaetatga	19
S2108 3411	
H211H 19	
HOICH DNA	
<pre>%2130 Homo sapiens</pre>	
(22 ti)	
<2.30 Cyclin Bl ribozyme binding site	
4400% 3411	

- '

tgttggasta tgacatggt	19
+0010 + 3410 +0010 + 19 +0010 + DNA +0013 + Homo sapiens	
+2220 + + Cyclin B1 ribbzyme binding site	
<pre>-0400 * 3412 tggtgcaptt tectootte</pre>	19
<pre>#210 + 3413 #211 + 19 #212 ** DNA #213 # Homo sapiens</pre>	
<pre>&lt;220 - &lt;223 - Cyclin B1 ribozyme binding site</pre>	
<400. 3413 ggtgeachtt colocttot	19
<pre>&lt;2100-3414 &lt;2210-19 &lt;2210-EMA &lt;22130-Homo sapiens</pre>	
<pre>&lt;2200 &lt;22100 &lt;22100</pre>	
<400> 3414 qtqqaactite otoottoto	19
+02100+ 3415 +02110- 19 +02120- DNA +02130- Homo sapiens	
-07.700- -07.730- Cyclin B1 ribozyme binding site	
+:4008 3415 dactitecto etteteaaa	19
+0100 3416 +0110 10 +0210 DNA +0213> Homo sapiens	

<pre><pre><id20 *="" b1="" binding="" cyclin="" pre="" ribozyme="" site<=""></id20></pre></pre>	
0400 · 3416 intootoott otcaaattg	19
H210 + 3417 H211 + 19 H211 + DNA H213 > Homo sapiens	
<pre>0220 + 0223 * Cyclin B1 ribozyme binding site</pre>	
0400 × 3417 ttootootto toaaattgo	19
<pre>&lt;210 &gt; 3418 &lt;2210 &gt; 19 &lt;210 &gt; DNA &lt;210 &gt; Homo sapiens</pre>	
+220 + +223 + Cyclin B1 ribozyme binding site	
<pre>&lt;:400 + 3418 cotcottoto aaattgcag</pre>	19
<pre>0010 - 3419 0011 - 19 0010 - DNA 0013 - Homo sapiens</pre>	
-0200	
-04000-3419 ttotoasatt goagbagga	19
+00100+ 3420 +0010+ 19 +0010+ DNA +00138 Homo sapiens	
HCCLOCK HCCCOCK Cyclin B1 ribozyme binding site	
-(400)- 3420 gcaggagott tttgcttag	19
-:210:- 3421	

HD1C → DNA	
(213 · Homo sapiens	
::200 ×	
(1.3) « Cyplin B1 ribbzyme binding site	
	1:
daggagottt tigottago	1
<210 + 3422	
(211 - 19)	
d110 ← DMA	
CO13> Homo sapiens	
<220>	
<223> Cyclin B1 ribozyme binding site	
<400 > 3422	
aggagotitt tgottagoa	1:
<210 × 3423	
<211 × 19	
H212 > DNA	
313> Homo sapiens	
C229A	
22230 Cyclin B1 ribozyme binding site	
Type Direction of the state of	
(400) - 3423	
qqaqettttt gettageae	19
22.50	
<pre>&lt;0.100 34.24 </pre>	
(211): 19	
G2129-DMA	
(1130 Homo sapiens	
(220)	
Class Cyclin El ribozyme binding site	
4000-3424	
mittitgett agcaetgaa	1 9
C10:- 3425	
12110-19	
1211: DNA	
121:00 Homo sapiens	
(226)	
(225) Cyclin El ribozyme binding site	

.

<400 * 3425	
tottigotta gcactgaaa	19
H210 + 3426 H211 + 19 H212 + DNA H313 + Homo sapiens	
<pre><pre><pre></pre> <pre><pre></pre> <pre><pre></pre> <pre>C220 &gt;</pre> <pre></pre> <pre><pre></pre> <pre></pre> <pre></pre></pre></pre></pre></pre></pre>	
<pre>&lt;400 x 3426 actgaaaatt ctggataat</pre>	19
H210 + 3427 H211 > 19 H212 > DNA H213 > Hamo sapiens	
<pre>&lt;2208 &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
<pre>&lt;400&gt; 3427 otgaaaatto tggataatg</pre>	19
<pre>&lt;210&gt; 3428 &lt;2211&gt; 19 &lt;2212&gt; DNA &lt;2213&gt; Homo sapiens</pre>	
<2208 <2238 Cyclin B1 ribozyme binding site	
-:4005 3428 attotggata atggtgaat	19
HO10% 3409 HO11% 19 HO110% DNA HO110% Homo sapiens	
<pre>-direc</pre>	
+04000-3409 acadeaacto tacaacatt	19
+00100+ 3430 +00110+ 13 +001.0+ DNA +00130+ Homo sapiens	

+:220+ +:223+ Cyclin Bl ribozyme binding site	
+(400 + 3430 appaactota caacattac	19
+1010 + 3431 +1011 + 19	
HOBIO - DNA HOMO sapiens	
<pre>&lt;:220 - &lt;:223 - Cyclin B1 ribozyme binding site</pre>	
+:400+ 3431 otabaabatt acotgtoat	19
<pre>#210&gt; 3432 +211&gt; 19</pre>	
<pre>#212 * DNA #213 * Home sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Cyclin B1 ribozyme binding site</pre>	
+:400 × 343.	
tacaacatta cotgecata	19
H210H 3433	
+1211× 19	
HC2125 DNA HC2135 Hcmc sapiens	
-undo-s	
-MLDB: Cyclin B1 ribozyme binding site	
s:400.5 3433	
attacotyto atatactya	19
+:210:- 3434	
(211): 19	
+02125 DNA +02135 Homo sapiens	
ALLO DAME Superior	
-02200-	
+21150 Cyclin B1 ribozyme binding site	
·:400:- 3434	
acctgtcata tactgaaga	19

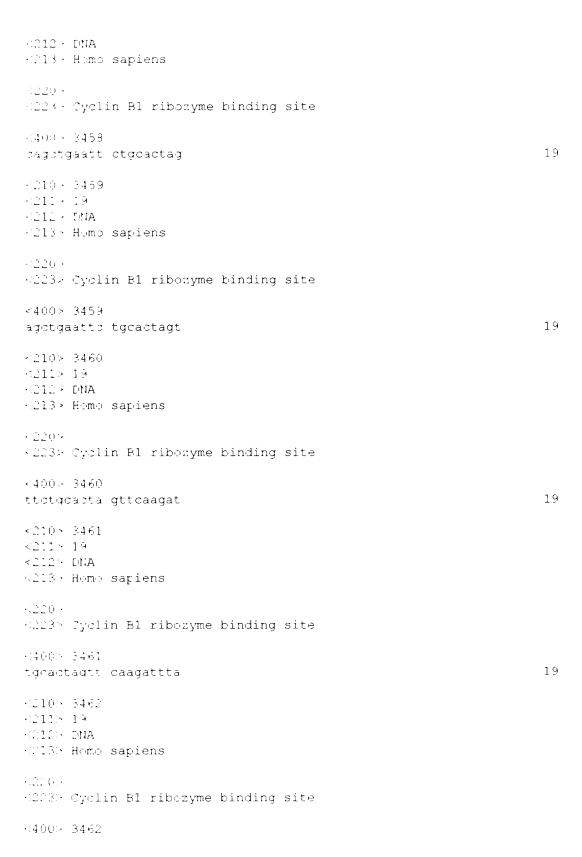
H210+3435 H211+19 H113+ Homo sapiens	
+220 + +2213 + Cyclin Bl ribozyme binding site	
-:400 - 3435 Otgtoatata otgaagaat	19
+1010 + 3436 +1011 + 19 +1011 + DNA +1013 + Homo sapiens	
<pre>k220. <pre>c223.3 Cyclin B1 ribozyme binding site</pre></pre>	
<400 - 3436 Stgaagaate tettettee	19
<pre>&lt;210 * 3437 +2211 &gt; 19 &lt;212 &gt; DNA &lt;213 &gt; Homo sapiens</pre>	
<pre>&lt;:220 * -:223</pre>	
+:400> 3437 gaagaatete ttetteeag	19
+0010% 3438 +0011% 19 +0010% DNA +0013% Homo sapiens	
+Clous +Clous Cyclin B1 ribozyme binding site	
+:4005 3438 agaatotott ottopagtt	19
+C2108 3439 +C2118 19 +C2118 DNA +CC188 Homo sapiens	
ROMECO: RECORD Cyclin El ribozyme binding site	

<pre>&lt;4)0+ 3439 quatototto ttocagtta</pre>	19
+1210 + 3440 +1211 + 19 +1212 + DMA +1213 + Homo sapiens	
+220 + +223 + Cyclin B1 ribozyme binding site	
+400+ 3440 atotottott coagttatg	19
+210 > 3441 +211 > 19 +212 > DNA +213 > Homo sapiens	
+00008 +0003> Gydlin B1 ribozyme binding site	
+:400 × 3441 totottotto cagttatgo	19
<pre>%U10 + 3442 %U10 + DNA %U10 * DNA %U13 * Homo sapiens</pre>	
-0008 -0008 Cyclin Bl ribozyme binding site	
04005 3442 tottocadtt atgoagoac	19
+00108 3443 +00108 19 +00108 DMA +00188 Homo sapiens	
+:27.0- +:27.3- Cyclin B1 ribozyme binding site	
<pre>+:400+ 3443 ontocag*ta tgcagcacc</pre>	19
+001()+ 3444 +001()+ 19 +001()> FNA	

<pre><c130 homo="" pre="" sapiens<=""></c130></pre>	
+020+ +023+ Gyolin B1 ribozyme binding site	
Systin bi iitoszyme binding site	
$\pm (400 \pm 3444)$	
dapatggata agaatgtag	19
H210 + 3445	
H211 × 19	
H212 + DNA	
+213 · Homo sapiens	
N220 +	
<223 - Cyclin B1 ribozyme binding site	
4400 + 3445	
taaqaatgta gtoatggta	19
+210 + 3446	
H211 + 19	
<pre><dim> DNA</dim></pre>	
<pre>&lt;213 · Homo sapiens</pre>	
+:220%	
+223: Syclin B1 ribozyme binding site	
+:400:5-3446	
gaatgtagte atggtaaat	19
+2108 3447	
K2115-19	
ROTES DNA	
+213+ Homo sapiens	
+:::2:::::::::::::::::::::::::::::::::	
-12.13 - Cyclin El ribozyme binding site	
+14000+ 3447	3.0
agticatgota aatcaagga	19
+:21:0:+ 3448	
R2119 19	
SCIID ONA	
<pre>+0.130 Homo sapiens</pre>	
- (III 0)-	
R2030 Cyclin Bl ribozyme binding site	
<4000 3448	
atggtaaatc aaggactta	19

-1210 →	3.44.9	
<1.211 ×		
1		
	Homo sapiens	
-1220 -		
S.1.1.3 *	Cyclin B1 ribozyme binding site	
4(4())) 두	2146	
		19
ं अववपुर्	gaett acaaageac	15
-(210) -	3.450	
<211>		
<2112>		
\3>	Homo sapiens	
<220>		
	Cyclin B1 ribozyme binding site	
~/	cyclin bi libozyme binding site	
<400>	3.450	
	actta caaagcaca	19
	10 50 11 Cadagouca	
s210>	3451	
<211>		
<212>		
	Homo sapiens	
Security 1	Tables Suprems	
<0.00>		
S. M. 13 5 5	Cyclin B1 ribozyme binding site	
<400>	3451	
cat.gac	stats aagaacaag	19
<210×	3452	
. 111	19	
<	ZNA	
42135	Homo sapiens	
4111 (CS		
<12.230×	Cyclin E1 ribozyme binding site	
·[40]	3452	
aqaaca	aadta tgccacatc	19
·2100		
-12111-		
.12171-		
.12150	Homo sapiens	
·12201-		

-223 - Oyolin Bl ribozyme binding site	
·(40) + 3453	
at posacato gaagcatgo	19
<2.210 + 3454	
42.11 + 19	
<212 + EMA	
<pre>&lt;213 * Home sapiens</pre>	
-220 -	
+323 + Cyclin Bl ribozyme binding site	
-:400 +: 3454	
aagbatgota agatbagba	19
<210 × 3455	
<2115-19	
KIII- DNA	
<2135 Homo sapiens	
<220.8	
<pre><!--D23% Cyclin B1 ribozyme binding site</pre--></pre>	
<4005-3455	
tgotaagato agcactota	19
+2105-3456	
H211N 19	
<pre>Home sapiens</pre>	
-: <u>0.00</u> 0-	
<pre></pre> <pre></pre> <pre>C2238 Cyclin B1 ribozyme binding site</pre>	
(400): 3456	
atcaqcactc taccacag:	19
HD109-3457	
-C115-19	
HONON DNA	
:C13: Home sapiens	
H2008	
HIDAM Cyclin Bl ribezyme binding site	
-:4c0:- 3457	
dagdactdta ccacagetg	19
42110-19	
· was a second of the second o	



gcantagtto aagatttag	19
<210 ← 3463	
4011 - 19	
HOLLS DNA	
<pre>#C13 * Homo sapiens</pre>	
<ul> <li>€1.12.0 +</li> </ul>	
+22% + Cyclin Bl ribozyme binding site	
<i400 +="" 3463<="" td=""><td></td></i400>	
gttcaagatt tagecaagg	19
<210 - 3464	
<211 - 19	
<pre></pre>	
<213 · Homo sapiens	
<0.00 •	
<0003 Cyclin B1 ribozyme binding site	
<400 > 3464	
ttbaagattt agccaaggc	19
<0.10 > 3465	
<2115-19	
ROLLY DNA	
<pre>&lt;:Cl3 + Homo sapiens</pre>	
<020 -	
K223 · Cyclin B1 ribozyme binding site	
<400 \ 3465	
toaaqattta godaaggot	19
-:210 - 3466	
4011 + 19	
KO11. DNA	
-113. Homo sapiens	
ricaer-	
-DD33- Cyclin B1 ribozyme binding site	
-:400:- 3466	
daaaggtgta acttgtaaa	19
-C10-3467	
HIII: 19	
H21LH DNA	
:2152 Homo sapiens	

<pre>::220 * ::::3 * Cyclin B1 ribozyme binding site</pre>	
H400+ 3467 ggtqtaastt gtaaacttg	19
+1210 + 3468 +1211 + 19 +1212 + DNA	
#213 + Homo sapiens	
<pre>edd0. edd0. e</pre>	
<400 + 3468 gtaacttgta aacttgagt	19
<pre>&lt;210&gt; 3469 &lt;211&gt; 19 &lt;212&gt; DNA &lt;213&gt; Homo sapiens</pre>	
-MANA Cyclin B1 ribozyme binding site	
<pre></pre> <pre>-(400&gt; 3469 ttqtaaactt gagttggag</pre>	19
<pre>&lt;0.10 + 3470 &lt;0.11 + 19 &lt;0.10 + 5MA &lt;0.13 + Homo sapiens</pre>	
<220%	
<pre><c23> Cyclin B1 ribozyme binding site</c23></pre>	
<pre>0400 + 3470 aacttgagtt ggagtacta</pre>	19
+0010 × 3471 +0211 × 10 +0210 × 10A +0213 × Homo sapiens	
eggans	
-0.23 Cyclin B1 ribozyme binding site	
::400:- 3471 adt:ggagta ctatacttt	19
K210% 3472	

<pre>42:11 + 19 42:12 + DMA 42:13 + Homo sapiens</pre>	
<pre><pre><id20 +="" <pre=""></id20></pre> <pre><id23 *="" b1="" binding="" cyclin="" pre="" ribozyme="" site<=""></id23></pre></pre>	
<pre>&lt;400 - 3472 typagtabta tactttaca</pre>	19
<pre>*D10 &gt; 3473  <d11> 19  <d11> DNA  *D13 &gt; Homo sapiens</d11></d11></pre>	
<pre><dd0> <d133> Cyclin B1 ribozyme binding site</d133></dd0></pre>	
<400 > 3473 gagtactata ctttacaaa	19
<pre>\$210 &gt; 3474  *2211 &gt; 19  \$211 &gt; DNA  \$213 &gt; Homo sapiens</pre>	
<pre>&lt;:220 + &lt;:223 &gt; Cyclin B1 ribozyme binding site</pre>	
+4005 3474 tactatactt tacaaacta	15
K210	
+:2100+ +:2130+ Cyclin B1 ribozyme binding site	
-:400:- 3475 actatacttt acaaactaa	19
<pre>sm10s 3476 +m11s 19 +m12:- DNA +m12:- Homo sapiens</pre>	
<pre></pre> <pre>&lt;</pre>	

-(400) - 3476 -ctatactita caaactaaa	19
<pre>%210 + 5477 %211 + 19 %212 + DNA %213 + Homo sapiens</pre>	
<pre>-::::0 + &lt;::::::::::::::::::::::::::::::::::::</pre>	
k400 + 3477 ttabaaasta aaattggca	19
<pre>K210 + 347% K211 + 19 K213 + DNA &lt;213 + Homo sapiens</pre>	
<pre>&lt;22008 &lt;2223   Cyclin B1 ribozyme binding site</pre>	
H400+ 3478 aactaaaatt ggcacatgt	19
<pre>K010 % 3479 H0011 + 19 H011   EMA + 213   Homo sapiens</pre>	
<2208 <2238 Odc15 hs ribozyme binding site	
H400 - 3479 aggaagssto tgagtooga	19
+1210 + 3480 +1211 + 19 +1212 + DNA +1213 + Bomo sapiens	
+02200+ +02235 Cdc25 hs ribozyme binding site	
+:400+ 3480 actotgagto ogaogtigg	19
<pre>%2100 3481 %2110 14 %2170 DNA %2170 Home sapiens</pre>	

<pre><!--220 + -:223 - Odc.15 hs ribozyme binding site</pre--></pre>	
(400 + 3481 gtoogacytt ggeetaees	19
+010 × 3480 +011 × 19 +010 × DNA	
*213 * Homo sapiens	
<pre>k:220 + :223 &lt; Ode25 hs ribezyme binding site</pre>	
<pre>-:400 + 3482 ogttggodta cocagtogg</pre>	19
<pre>&lt;210 &gt; 3483 &lt;2211 &gt; 19 &lt;0123</pre>	
<pre>&lt;020s &lt;223&gt; Odo05 hs riboxyme binding site</pre>	
k400% 3483 otacocagto ggaaggoag	19
<pre> <pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>0.00s -0.03s Ode05 hs ribozyme binding site</pre>	
<400% 3484 agotqcaato tagttaact	19
KC10% 348% KC11% 19	
+00120+ DNA +00130+ Home sapiens	
<pre>+DD00: +DD03: Cdc.15 hs ribezyme binding site</pre>	
<14002 3485 otgoaatota gttaactac	19

H010 > 3485 H011 + 19 H012 + DNA H0m0 sapiens	
+1230 × +1230 + Chol5 hs ribozyme binding site	
-:400 - 3486 -caatetagtt aactacete	19
+210 + 3487 +211 + 19 +212 > DNA +213 + Homo sapiens	
<pre>&lt;220x &lt;223&gt; Cdc25 hs ribozyme binding site</pre>	
<400> 3487 aatotagtta actacotoo	19
<pre>&lt;010 &gt; 3488 &lt;0211 &gt; 19 &lt;0212 &gt; DNA &lt;0213 &gt; Homo sapiens</pre>	
-inno- -inno- Odono hs ribozyme binding site	
<pre>+400/2 3488 tagttaacta cotoottto</pre>	19
<pre>&lt;:2108 3489 &lt;:2118 19 &lt;:2128 DNA &lt;:2138 Himo sapiens</pre>	
-mags -mags Cdc25 hs ribozyme binding site	
-:400:- 3489 taactacato otttoccot	19
+U10: 3490 +U11: 19 +U11: DNA +U12: Homo sapiens	
+12C> +123> Cdc25 hs ribozyme binding site	

+:400 + 3490 otapetestt tedectaga	19
<pre>8210 + 3491 -2211 + 19 -2212 + DNA -2213 + Homo sapiens</pre>	
HCCCO + CMcCCO hs ribozyme binding site	
+(400 + 3491 tabataatti adadtagat	19
+1210 x 3492 +1211 x 19 +1212 x DNA +1213 x Homo sapiens	
<pre></pre> <pre>&lt;220&gt; &lt;223&gt; Odc25 hs ribozyme binding site </pre>	
<pre>&lt;400% 3492 abstraction contagatt</pre>	19
+C210 > 3493 +C211 > 19 +C210 > CNA +C213 > Homo sapiens	
<pre>&lt;220&gt; &lt;223 Odc.5 hs ribozyme binding site</pre>	
<pre>&lt;:400 + 3493 ettteesta gattteett</pre>	19
+210× 3494 +2115 19 +2113× DMA +2113× Homo sapiens	
<pre>kmi0s &lt;12.0s &lt;12.3s Odd05 hs ribozyme binding site</pre>	
+400+3494  cocctagat1 tcctttcat	19
+:210:+ 3495 +:211:+ 1 ) +:212:- FNA	

-:213 · Homo sapiens	
+1239 +	
$\pm 223 \pm \text{Cdc25}$ hs ribozyme binding site	
÷:4au + 3495	
Destagatt cottoatt	19
+1110 + 3496	
HILLS 19 HILLS DNA	
H213 Homo sapiens	
<pre>&lt;220 - 222</pre>	
$\pm 223 \pm 0$ dd25 hs ribozyme binding site	
<400 × 3496	
octagatito otticatio	19
+:210 × 3497	
4211 + 19	
RICTO + DNA	
-213 Homo sapiens	
<220 S	
<pre>&lt;223&gt; @dc25 hs ribozyme binding site</pre>	
+:400% 3497	
agatttoott toattotgo	19
+2210 + 3498	
+:211> 19 +:212> DNA	
+213: Homo sapiens	
+:2109 +:2138 Cdc25 hs ribozyme binding site	
The state of the s	
+:4008 3498	
gatticettt cattetget	19
HD10H 349H	
02118 19	
<pre></pre>	
+212+ Homo sapiens	
40.20h	
<pre><!--2230 Cdc05 hs ribozyme binding site</pre--></pre>	
-(40D)- 349⊕	
atttocttte attetgete	19

+:010+ 3500 +::11+ 19 +::12+ DMA +::13+ Homo sapiens	
<pre>&lt;220 + &lt;223 - Cdc25 hs ribbzyme binding site</pre>	
-:400 - 3500 toctitoatt otgotoaag	19
<pre>40.10 + 3501 4011</pre>	
<220> <223> Odc25 hs ribozyme binding site	
<pre>&lt;400&gt; 3501 cotttcattc tgctcaagt</pre>	19
<pre>&lt;010&gt; 3502 &lt;011&gt; 19 &lt;010&gt; DNA &lt;013&gt; Romo sapiens</pre>	
-CD098 -CD03> Cdc05 hs ribozyme binding site	
k400> 3500 pattotgoto aagtottog	19
+0.10 × 3503 +0.11 + 19 +0.10 × DNA +0.13 × Homo sapiens	
-2260- -2230- Cdc.5 hs ribozyme binding site	
-4005 3503 tgotdaagte ttogootgt	19
*0100+ 3504 *0110+ 14 *0100+ DNA +0130+ Homo sapiens	

4220>

<pre><!--223</th--><th></th></pre>	
-:400 + 3504 blosagtoti ogcotgtgt	19
+010 + 3505 +011 + 19 +012 + DNA	
+213 + Hom → sapiens	
+20n+	
<pre>silil3 - Odcl5 hs ribozyme binding site</pre>	
k400 + 3505	
toaaqtotto gootgtgto	19
<010 ≠ 3506 <011 ÷ 19	
KLTT: T9 KLTL: DNA	
<213 · Homo sapiens	
<220 ×	
-2235 Odc25 hs ribozyme binding site	
<400 \( \text{3506} \)	1.0
ogootgtqto ogatoocta	19
K210 + 3507 K211 + 19	
KOION DNA	
S213 Home sapiens	
40 NG 8	
-1.13 + 0.dc.15 hs ribozyme binding site	
-:400 -: 3507	1.0
gtgtbogatb octatotac	19
H210 × 3509	
CC11 + 19	
OFINA DNA OFINA Homo sapiens	
<pre>%2000 - %2003 - Odc25 hs ribozyme binding site</pre>	
(400) 3508	
ooga: coota totacttto	19
<210:- 3509	
K2111/- 19	

3213 - DNA	
-0013 - Homo sariens	
F1129 -	
FRANK - CHall his ribozyme binding site	
11.00	
H400 + 3500	
yatoodtato tadtitoto	19
HIII 0 + 3510	
+1011 + 19	
HILL HIDNA	
+1113 + Homo sapiens	
The time baptone	
s(0,00 ×	
<pre><dm3> Odc25 hs ribozyme binding site</dm3></pre>	
4400 0140	
<4400 × 3510	
todetateta ettitetete	19
+:210 + 3511	
K211 × 19	
H212 M DMA	
S213 Homo sapiens	
Description of the second of t	
+122205	
+ 223 + Cdc15 hs riboxyme binding site	
1100 2511	
+1400+13511	
statotastt tototooto	19
+IZ10> 351I	
H211> 19	
FOLOS EMA	
Hama sapiens	
and all	
+:210:4	
<22330 Cdc25 hs riboxyme binding site	
+:400:+ 351.1	
	2.0
tatotaciti ototootot	19
+00100-3513	
+0.115 19	
AUCCOLOUS AND STATE OF THE STAT	
H2130 Homo sapiens	
+(210)	
-2130 Cdcl5 hs ribozyme binding site	
+:4002 3513	
**VU/	

atstacttts teteetett	19
+1310 + 3514 +1311 + 14 +1312 + DNA +1313 + Homo sapiens	
+0.00 + +0.005 hs ribozyme binding site	
+(400) + 3°14 otaotttoto tootottgt	19
+1010 + 3515 +1011 + 19 +1012 + DNA +1013 + Homo sapiens	
+030> <223 - Odc15 hs ribozyme binding site	
<pre>&lt;400 + 3515 actitetete etettgtag</pre>	19
<pre>&lt;210% 3516 &lt;221% 19 &lt;221% DNA &lt;221% Home sapiens</pre>	
<pre><pre><pre><pre><pre></pre> <pre><pre><pre></pre> <pre></pre> <pre>Class Cdc25 hs ribozyme binding site</pre></pre></pre></pre></pre></pre></pre>	
r400% 3516 thatateata tigitageaa	19
+01108 3517 +0114 17 +0012 DMA +0115 Homo sapiens	
+MANON- +MANON Cdc25 hs ribozyme binding site	
+4400+ 3517 dididetett gtageaage	19
+0.100+ 3918 +0010+ 19 +0210+ DNA +0213> Homo sapiens	

+:220 · +:223 · 3dc25 hs ribozyme binding site	
+:400 + 3518 testettata geaageete	19
+0010 + 351 + + + + + + + + + + + + + + + + + + +	
Homo sapiens	
<pre>+:220 + +:213 + Ode5 hs ribozyme binding site</pre>	
<pre>&lt;400% 3519 aqcaagcotc agactccag</pre>	19
<pre>&lt;010 &gt; 3520 &lt;011 &gt; 19 &lt;011 &gt; 5NA &lt;0215 &gt; Homo sapiens</pre>	
<pre>&lt;:220&gt; &lt;220&gt; &lt;2223&gt; Cdc25 hs ribozyme binding site</pre>	
<pre>+:400% 3520 deteagaete caggettga</pre>	19
*010% 3501 *011% 19 *011% DNA	
<pre>SD13&gt; Homo sapiens</pre>	
+:22%+ Odd25 hs ribozyme binding site	
+400+ 3501 otocaggott gagotaggt	19
KO100-3500 -0010-19 -0010-DNA	
SC130 Homo sapiens	
+2100+ +2103+ Odc25 hs ribozyme binding site	
equin 3511 qui dageta ggittigit	19
H2102 3523	

+CC11+ 19 +CC12+ DNA +CC13+ Homo sapiens	
<pre>&lt;1200 - &lt;1203 - Odc05 hs ribozyme binding site</pre>	
<pre>-:400 - 3523 yayatagytt ttgtttttc</pre>	19
<pre>0210 + 3524 0211 + 19 0212 + DMA 0213 + Homo sapiens</pre>	
<pre>&lt;2200p &lt;2210p &lt;2213p Odc25 hs ribozyme binding site</pre>	
<pre>&lt;400&gt; 3524 agotaggtit tgttttct</pre>	19
<pre>&lt;210&gt; 3525 &lt;2211&gt; 19 &lt;2212&gt; DNA &lt;2213&gt; Homo sapiens</pre>	
<pre>Algonia Algonia A</pre>	
<pre>-:400 % 3525 getaggtint gttttete</pre>	19
+010× 3506 +0011× 19 +0010× DNA +0013× Homo sapiens	
HDDGG Cddl5 hs ribozyme binding site	
H400% 3526 addititiant ittleteetg	19
+00100+ 3527 +0010+ 19 +0010+ DMA +0010+ Homo sapiens	
+0.200+ +2200+ Cdc25 hs ribozyme binding site	

H400 + 3527 gattitgitt ttotootgg	19
H010 + 3508 H011 + 19 H010 + DMA	
40713 - Homo sapiens	
<pre>&lt;120 * &lt;123 * Odo15 hs ribozyme binding site</pre>	
स्थ्ये । 3528 प्राचित्रप्रदेशी telectggt	19
<pre>&lt;010 3529 &lt;011.&gt; 19 &lt;010 3929</pre>	
<pre>cdld&gt; DNA kdld&gt; Homo sapiens</pre>	
<pre>&lt;0000 &gt; &lt;0000 f</pre>	
<400 × 3529 tittgttttt ctcctggtg	19
+010 × 3530 +011 + 19 +010 + 5NA	
+2213c Homo sapiens	
<pre><datable <="" pre=""> <pre><datable <="" pre=""> <pre><datable <="" pre=""> <pre><datable <="" pre=""> <pre></pre> <pre>&lt;</pre></datable></pre></datable></pre></datable></pre></datable></pre>	
N400N 3530 titattitin teetggtga	19
+0100+ 3531 +0110+ 19 +0100+ INA	
+Cli30 Homo sapiens	
+C220:- +C230- Cdc25 hs ribozyme binding site	
+44000 3831	10
tqtf:ttctc ctggtgaga	19
+021(0) 3002 +0211: 13	
HILLS DNA	
<2130 Homo sapiens	

<pre><dd:: <pre="">Cdc25 hs ribozyme binding site</dd::></pre>	
H400+ 3532 ggtgagaatt ogaagacca	19
R210 + 3533 R211 + 19	
HOBIER DWA HOMO sapiens	
<pre>+MAGO+ +MAGO+ CdcA5 hs ribozyme binding site</pre>	
+:400× 3533 gtgagaatto gaagaccat	19
+:210 × 3534 +:211 × 19	
<pre>&lt;012&gt; ENA &lt;013&gt; Homo sapiens</pre>	
<pre>%2208 %2238 Odc25 hs ribczyme binding site</pre>	
-0400 × 3534 agabbatgto taoggaabt	19
<pre></pre> <pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><p< td=""><td></td></p<></pre>	
<210> DNA <213> Homo sapiens	
kmman kmman kmmman	
<pre>&lt;40ub 3535 accatgteta eggaactet</pre>	19
KD105 3536 KD15 19	
<pre>%Alto Ta <pre>%Did&gt; DNA <pre>%Did&gt; Homo sapiens</pre></pre></pre>	
<pre><i210> <i223> Cdc.5 hs ribozyme binding site</i223></i210></pre>	
<4400° 3536	
tacggaactc ttctcatcc	19

HC10 + 3537 HC11 + 10 HC12 + DNA HC13 + Homo sapiens	
<pre>%2208 %223 * Odo25 hs ribozyme binding site</pre>	
0400 · 3537 σημααρτοτί otcatocae	19
0010 - 3538 0011 - 19 00102 DNA 0013 - Home sapiens	
<220> <223> Odo25 hs ribozyme binding site	
<400> 3538 ggaactotto toatocaba	19
<pre>&lt;2108 3539 +2118 19 &lt;2128 DNA +2138 Homo sapiens</pre>	
HUDO: HUD3 + OdoN5 hs ribozyme binding site	
-:400. 3539 aactottoto atocacaag	19
HD10:- 3540 HD11:- 19 HD12:- DNA HD13:- Homo sapiens	
+Times +Iii30+ Cdcl5 hs ribozyme binding site	
<pre>+:4000- 3540 tottotoato cacaagaga</pre>	19
+02100+ 3541 +0210+ 19 +0212+ DMA +0212+ Homo sapiens	
<pre>&lt;22.00 &lt;22.30 Cdc25 hs ribozyme binding site</pre>	

4213 · Homo sapiens	
<pre>%200 + %200 + Odo05 hs ribozyme binding site</pre>	
องสอบ - 3546 องเวลซุปประวัติ ggtotaato	19
+010 + 354° +0110 19 +0100 DNA +013° Homo sapiens	
<pre>&lt;220* &lt;223* 0dm25 hs ribozyme binding site</pre>	
k400> 3547 gttttaggtc taacaaagg	19
<pre>&lt;210&gt; 3548 &lt;211 * 19 &lt;210 * DNA &lt;213 * Homo sapiens</pre>	
<220> <220> <223+ Odd25 hs ribozyme binding site	
<400 - 3548 tttaggtota atcaaagga	19
<pre>SD10 + 3549 SD11 &gt; 19 <d12> DNA +D13 + Homo sapiens</d12></pre>	
<pre>%220 * %220 * Cdc25 hs ribozyme binding site</pre>	
ार्बिक्तार 3549 angtotaato aaaggaaaa	19
<pre><cli>Clic 3550 <clic 19="" <clic="" dna="" homo="" pre="" sapiens<=""></clic></cli></pre>	
+00000+ +00000+ Cdc05 hs ribozyme binding site	
+:400% 3550	19

<pre>H210 + 3551 H211 + 10 H212 + DMA H213 + Homo sapiens</pre>	
-020 - -020 - 34c25 hs ribozyme binding site	
+:400 + 3551 qaaaatgtta aacctgoto	19
+:210 + 3552 +:211 + 19 <:212 + DNA +:213 + Homo sapiens	
<pre>&lt;220 + &lt;223 + Odc25 hs ribozyme binding site</pre>	
<pre>&lt;400 + 3550 asacctgete otggagaga</pre>	19
+(210 + 3553 +(211 + 19 +(212 + DNA +(212 + Homo sapiens	
<pre>+220 +220 +220 +220 *Cdc15 hs ribozyme binding site</pre>	
<4000 3553 agagacact: octttaccg	19
+(210) 3854 +(211) 19 +(212) DNA +(213) Home sapiens	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>84000 3554 gagadacttc ctttaccgt</pre>	10
+02100+ 3856 +00110+ 19 +0.110+ DNA +00130+ Hemo sapiens	
+:220;z	

-

-:223 · Odc25 hs ribozyme binding site	
+400 + 3555 anabitobtt taccgtotg	19
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>%113 * Home sapiens</pre>	
<pre><pre>&lt;0.00 * Cdc25 hs ribozyme binding site</pre></pre>	
-:4003556 cactteettt accgtetgt	19
<010> 3557 <011> 19	
<pre>*Cl3&gt; DNA *Cl3&gt; Homo sapiens</pre>	
<22007 <2203   Odc25 hs ribozyme binding site	
<400% 3557 acttocttta cogtotgto	19
<0.10 × 3658 <0.11 × 19	
<pre><c11.5 <213="" dna="" homo="" pre="" sapiens<=""></c11.5></pre>	
<pre>&lt;210.+ &lt;223    Cdc25 hs ribozyme binding site</pre>	
-4000-3558 stitaccyts tgtocagat	19
+:216:+3559 +:211:+19 +:212:+DNA +:213:+Homo sapiens	
-CLTC). -CLTC: Cdc25 hs ribozyme binding site	
::400: 3°54 accgfctata cagatgtec	19
-:::10:- 3560 -::11> 14	

<pre>0212 + DNA 0213 + Homo sapiens</pre>	
<pre>&lt;2200 + &lt;2223 + Odol5 hs ribozyme binding site</pre>	
<pre>&lt;400 + 3560 topagatqto cotagaact</pre>	19
+0210 + 3561 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
+0000+ +0003+ Odc05 hs ribozyme binding site	
<pre>-400&gt; 3561 gatgtocota gaactccag</pre>	19
+310 × 3562 +311 × 19 +3212 × DNA +3213 × Homo sapiens	
+:220> -:223 > Cdc25 hs ribbzyme binding site	
-:400 × 3562 octagaacto cagtgggca	19
<pre>&lt;0108 3563 &lt;0118 19 +02129 DNA +02138 Home sapiens</pre>	
+:210.+ +:210 Cacl5 hs ribozyme binding site	
H400m 3563 Nggqdaaatt tottggtga	19
+:210:- 3564 +:211:- 1* +:212:- 5MA +:213:- Hemo sapiens	
+02000 +02000 Cdc25 hs ribozyme binding site	
400 ≥ 3564	

ggg:aaattt cttggtgat	19
+1210 × 3565	
H211 + 19	
HOID + DMA	
H113 · Homo sapiens	
> <u>17_</u> (0)	
HALES CodeA5 hs ribozyme binding site	
+:400 + 3565	
ansaaattta ttggtgatt	19
a asaac 115 ccgg cgacc	1.7
KL109 3566	
<pre>&lt;211&gt; 19</pre>	
KO108 DNA	
*213> Homo sapiens	
<220>	
<pre><dm3> Cdc.15 hs ribozyme binding site</dm3></pre>	
<400 > 3566	
caaatttett ggtgattet	19
Samples ggegaeeee	10
<210> 3567	
K2118 19	
<pre></pre>	
+213× Homo sapiens	
s:220 s	
<pre>&lt;223 &gt; Odc25 hs ribozyme binding site</pre>	
+(4))) \cdot 356?	
stigatgatt otgoaaaco	19
+010x 3568	
<211 - 19	
HICTOR DNA	
+C2130 Homo sapiens	
+:2::0:+	
-:::3: Cdc::5 hs ribozyme binding site	
+:400+-3568	
tigatgatic tgcaaacct	19
K210:- 3569	
42117-19	
+C2170+ DNA	
<pre>42159 Homo sapiens</pre>	

-:22:4 -	
-0.23 · Odc. 5 hs ribozyme binding site	
<400 ← 3569	
typuaacota agcattttg	19
H119 + 3579	
HU11 + 19	
*1212 * DNA	
+1213 · Homo sapiens	
.sea	
+1013 + 3dc.15 hs ribozyme binding site	
s:400 × 3570	
	19
octaageatt ttgtotgga	15
<010> 3571	
<1.112 19	
K212S DNA	
<pre>8213@ Homo sapiens</pre>	
- Live Hall Suprone	
+:22a s	
<223 / Odc.15 hs ribozyme binding site	
<400 > 3571	
staagcatt. tgtotggag	19
+:210 × 3572	
<2211 × 19	
<pre>&lt;212 &gt; DNA</pre>	
<pre>&lt;213&gt; Homo sapiens</pre>	
+:220:+	
+0023+ Odc05 hs ribbzyme binding site	
(4.0 f) + 5 f 72	1 C
taaqcatttt gtotggagg	19
+1:10+ 3573	
HIGHTY INA	
+2130 Homo sapiens	
The second section is a second	
-223: Cac. 5 hs ribozyme binding site	
+:400:+ 3575	
qca: ttqtc tggaggaac	19
+21m+ 3574	

<211 - 19	
H212 + DNA	
<pre></pre>	
•	
×220 ×	
<pre></pre>	
-:401 - 3574	
ocaaaatgtt gootogato	19
222 Tude 400 good og atte	
H210 + 3575	
S211 + 19	
H211 - DNA	
+213 - Homo sapiens	
- 215 thanks Sapiens	
<2207	
<223% Cdc25 hs ribozyme binding site	
Kliba Cacia na Fibozyme binding site	
<400×3575	
angtigeste gatotites	19
anguigues gauduurug	1)
<210> 3576	
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<2112 DNA	
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K2208	
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K400× 3576	
tgostogato titogaato	19
eybblogacs cocogaate	1.0
KD105 3577	
<u118 19<="" td=""><td></td></u118>	
SULLE DNA	
<pre>%1135 Home sapiens</pre>	
+:000x	
+213> Odc15 hs ribozyme binding site	
*2237 CAGES IS FIDOZYME DINGING SICE	
+(400)+ 3577	
	19
detequatett togaatett	10
41.10% 367%	
60.00 to 50 to 60 to	
KITIDA DNA	
4.1130 Homo sapiens	
- 1900-	
-:220:-	
-223> Cdc25 hs ribozyme binding site	

0400 + 3573 Stagatottt ogaatotta	19
#210 + 3579 #211 + 19 #212 + DNA #213 + Homo sapiens	
<pre>H220 + H223 + Odo?5 hs ribozyme binding site</pre>	
<pre><ul><li><pre></pre></li><li><pre>togatettte gaatettag</pre></li></ul></pre>	19
<pre>SCD10 * 3580 CD11 * 19 CD10 * DNA <cd13 *="" homo="" pre="" sapiens<=""></cd13></pre>	
<pre><d20> <d20> <d203 *="" binding="" cdc25="" hs="" pre="" ribozyme="" site<=""></d203></d20></d20></pre>	
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<pre>KD10x 3581 KD11x 19 KD10x DMA KD13x Home sapiens</pre>	
<pre><i220s <="" pre=""> <pre><i223s binding="" hs="" odd25="" pre="" ribozyme="" site<=""></i223s></pre></i220s></pre>	
Hidus - 3581 ticquatott agcagtggg	19
+0110 + 3581 +0110 + 19 +0110 DMA +0110 Homo sapiens	
<pre>0.200 -0.200 -0.230 Odc.5 hs ribozyme binding site</pre>	
H400+ 348. togaatotta goagtgggg	19
+:21(:-3:8:- +:21(:-12 +::31:-DMA +:21::-Homo sapiens	

+:220 + +:223 + Cdc25 hs ribozyme binding site	
ा4000- 3583 tayagata actgccast	19
HC10H 3584	
H312 DNA H313 Homo sapiens	
<pre>+0.00.0 +0.00.25 hs ribbzyme binding site</pre>	
<400> 3584 actqccactc agcttacca	19
+:210 > 3585 +:211 > 19	
+C212> DNA +C213> Homo sapiens	
<pre>&lt;1208 &lt;12208 class Odc25 hs ribozyme binding site</pre>	
<pre><i400> 3585  bactcagett accaettet</i400></pre>	19
+210 + 3586 +211 + 19	
<pre><c113> DNA </c113></pre>	
+22008 +22238 Cdc25 hs ribozyme binding site	
<pre>&lt;400 &gt; 3586 actcagetta ccacttotg</pre>	19
+02108 3587 +02118 19	
<pre><m108 <m13="" dna=""> Homo sapiens</m108></pre>	
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+400 - 3587 cttaccactt ctgcagacc	19

<pre>KC10 + 3588 KC11 + 19 KC12 + DNA KC13 + Homo sapiens</pre>	
HI200. HI200. Odc25 hs ribozyme binding site	
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+0210 + 3589 +0211 + 19 +0212 + DNA +0213 + Homo sapiens	
<pre>&lt;220&gt; &lt;223 Gdc25 hs ribozyme binding site</pre>	
<pre></pre> <pre><pre>tqcaqacctt gatgaaact</pre></pre>	19
+1210 × 3590 +1211 + 19 +1212 + DNA +1213 + Home sapiens	
<pre>&lt;2200+ &lt;2230- Odol5 hs ribozyme binding site</pre>	
<pre>&lt;4000+ 3590 quauctgqtc acctggatt</pre>	19
+00103 3591 +00103 10 +00103 ENA +00103 Homo sapiens	
+02000 +00030 Odc25 hs ribozyme binding site	
+04000+ 3591 dacctggatt cttcaggac	19
+01100+3590 +0110+14 +0110+DMA +0110+Home sapiens	
<pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre>Cdc25 hs ribozyme binding site</pre>	

<pre>0400 * 3592 abstigation ticaggast</pre>	19
+1210 + 3593 +1211 + 19 +1211 + DNA +1213 + Homo sapiens	
+0.20 + +0.23 + Odo.25 hs ribozyme binding site	
-400 - 3593 - stygattott caggactto	19
<pre>%210 x 3594 %211 x 19 %212 &gt; DNA %213 x Homo sapiens</pre>	
<pre>0000 + 00003 Cdcl5 hs ribozyme binding site</pre>	
<pre>+(400% 3594) togattette aggaettea</pre>	19
+210 + 3898 +211 + 19 +212 + DMA +213 > Homo sapiens	
+:220 + +:223 + Cdc25 hs ribozyme binding site	
-:400 - 3595 ticangactt caggaagtg	19
+0210.+ 3+96 +0211.+ 14 +0210.+ ENA +0213.+ Homo sapiens	
+02205 +02205 Cdc15 hs ribozyme binding site	
+:40ut- 350b traddacttc aggaagtgc	19
+0110+3597 +0110+19 +2120+DNA	

+1213 + Homo sapiens	
+12.20 -	
+AAA3+ Odc25 hs ribozyme binding site	
±4.0.1 + 35.97	
waawigeatt tagetggga	19
H210 + 3598	
*12.11 + 19	
HILL HIMA	
+C13 / Homo sapiens	
*12.0 *	
+23:> 0dc25 hs ribozyme binding site	
<400× 3598	
aaqtgcattt acgtgggat	19
<pre>&lt;:010&gt; 3599</pre>	
40.115/19	
<pre>&lt;113&gt; 00A</pre>	
+C13> Homo sapiens	
<220%	
-0.03> Odc15 hs ribozyme binding site	
agtacattta gotgggatg	19
+(210× 3600	
+(211> 19	
HZ12> DNA	
<pre></pre>	
+:22as	
<pre>* &gt; Cia.5 hs ribczyme binding site</pre>	
+:4000× 3600	
gggatgaatc atgaccage	19
K2108 3601	
+211> 19	
-0.1128 DNA	
+0.1:5 Home sapiens	
+( <u>((), ())</u> +	
+270+ Cac25 hs ribozyme binding site	
<400% 3601	
ccagcaccta atgaaatgt	19

+:310 + 3602	
<0.11 - 19	
+0.12 + DNA	
<213 Homo sapiens	
all. Tom Suprens	
23.3m	
K020+	
RANAS Cirls hs ribozyme binding site	
<ul><li>&lt;1400 + 3602</li></ul>	
atgaaatgta goodagdad	19
S210 + 3603	
R211 + 19	
<pre></pre>	
<213 · Homo sapiens	
<220 /	
<223 Cdc25 hs ribozyme binding site	
<400 - 3603	
agpapagett etttgtage	19
aysasaysic occuysays	1.7
2310 S201	
<210 + 3604	
<211 - 19	
<212 - DNA	
<pre>k213 · Homo sapiens</pre>	
<220 -	
<pre>&lt;323 - Cdc25 hs ribozyme binding site</pre>	
The state of the s	
<400 + 3604	
	1.0
goadagetto tttgtagea	19
K210 + 3605	
<211,- 19	
KC10 - ENA	
<213 Homo sapiens	
•	
<1.00	
-:: Cdc. 5 hs ribozyme binding site	
The cast of its liberty me binding bite	
<4000 3605	
	1.0
acadottott tgtagcact	19
NI 100-3606	
+.111/+.14	
HIIIN DNA	
HO130 Homo sapiens	
rabie.	

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<pre>&lt;:223 • Odo25 hs ribozyme binding site</pre>	
0400 + 3606 payottottu gtagoapto	19
+0:10 + 360 / +0:11 + 10 +0:12 + DNA	
+:213 + Homo sapiens	
<pre><udlo-> <udlo-> &lt;</udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></udlo-></pre>	
ki400 / 3607 ottotttyta goadtooga	19
<210 + 3608 <211 + 19 <212 > DMA	
<pre>*D13* Homo sapiens</pre>	
<pre>&lt;0.20&gt; &lt;0.03* Odc25 hs ribozyme binding site</pre>	
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<pre>&lt;010x 3609 &lt;011x 19 </pre>	
<pre>&lt;0.12</pre>	
<pre>&lt;2208 &lt;2.38 Cdc.5 hs riboxyme binding site</pre>	
k400 - 3609 Dogaatgytt tygaccyty	19
+:110> 3610 +:11> 13	
<pre></pre>	
HILLONGO Cacl5 hs ribozyme binding site	
-(400)- 3/10 cdaatggftt ggaccgtgg	19
-0.10% 3/11 -0.210% 1/4	

-:212 - 1	DNA	
	Humu sapiens	
-1220 -		
	Odo25 hs ribozyme binding site	
·(4:)() ·	3611	
		19
. 1 . 5 5 5	3.3.2 9.4.4 9.4.9	
3.10 ·	วิที่ไ	
211		
-:212 · :		
	Homo sapiens	
	TONE - Supreme	
K220>		
	Odol5 hs ribozyme binding site	
1.2237 1	Jude 15 hs fibblyme binding site	
-:400× .	2012	
		19
geaalgi	tgta gttcatctg	19
2116.	5.41 %	
·0710 · .		
11>		
-0712 × 1		
HU137 1	Home sapiens	
-0.20%		
	Cdc. 5 hs ribezyme binding site	
	~	
-:4005		
aigtata	aqt: catctgcaa	19
41108 I		
4211 - 1		
40.12 × 1		
-0.1135 E	Homo sapiens	
-17.00		
	Cdc.5 hs ribozyme binding site	
·(400, - )		
tgtgtag	gito atotgoaaa	19
H2100-3		
47115		
·mili: I		
-12130 F	demo sapiens	
-(220)		
42230-0	Cdc25 hs ribozyme binding site	
<400 i − 3	3615	

gtaqttbatb tgbaaataa	19
<:210 x 3616	
+:211 + 13	
+:212 + DNA	
+:213 + Homo sapiens	
+:220 *	
+2223 + 0 4525 hs ribozyme binding site	
The Process of the Pr	
+:400 + 3616	
totquaaata aagaaaatg	19
+210 + 3617	
+1211 + 19	
+1212 + DNA	
+213 · Homo sapiens	
+:220×	
<pre></pre> <pre>&lt;</pre>	
LEGS (400.0 No Fired Minding Office	
<400 > 3617	
atqqaaactt ggtggacag	19
FD10F 3618	
+0112-19	
KO10 + INA	
+213 + Homo sapiens	
<2.20.*	
k223 - Cdc25 hs ribozyme binding site	
<400 · 3618	
awatmawata titgggcag	19
KD10:-3619	
(211 - 1)	
KI 100 TMA	
RC138 Bomo sapiens	
41.100	
-000000 hs ribozyme binding site	
(400) 361.4	
atgaaatati tgggcagtc	19
argadaca egggodgeo	+ -/
+2100+3620	
(0.11): 13	
-0.120- DNA	
KZ13> Homo sapiens	

Kaap •	
<pre>+CL23 + Cdc25 hs ribozyme binding site</pre>	
+(400 + 362 x	
tqaaatattt gggcagtoc	19
H210 + 3621	
+0.11 + 19	
SOLL SOLA	
<pre>&lt;:213 - Homo sapiens</pre>	
x:22a x	
+223 + Cds25 hs ribbzyme binding site	
+400 + 3621	
ttgggcagto ccattacta	19
K210× 3622	
×211> 19	
KD1D> DNA	
+013 Homo sapiens	
<2000×	
<pre>&lt;0.13&gt; Ode0.5 hs ribetyme binding site</pre>	
K400× 3600	
cadtoccatt actactgtt	19
k(210 k   360 k	
\$2119-19	
KONIN DNA	
xC135 Homo sapiens	
kd2us	
<pre><pre><pre></pre> <pre>Cdc.5 hs ribozyme binding site</pre></pre></pre>	
-:400× 3603	
aqthocatta ctactgttc	19
K210 x 3 x 2 4	
40.117- 19	
HILLICH DNA	
-1.150 Homo sapiens	
+0.200+	
-DDDS- Cdc25 hs ribozyme binding site	
-:40i::- 3n/4	
decattadta etgttecaa	19
#210% 3625	

·(211 · 1)		
<211 ← DNA		
-013 · Homo sapiens		
4220 ·		
<pre>x223 * Cdc25 hs ribozyme bind</pre>	ing site	
and cases no ribozyme bind	ing site	
-:400 + 3625		
	1	_
tadtactqtt ccaaaattg	1	9
<210 + 3626		
HIII + 19		
HIII2 + DNA		
<pre>Hill3: Homo sapiens</pre>		
<220>		
<pre>&lt;:213&gt; Cdc25 hs ribozyme bind.</pre>	ing site	
1		
<400 > 3606		
ttocaaaatt ggataaaaa	1	a
chocadate gyacadaaa	1	
<210> 3627		
<211> 19		
K212≻ DNA		
<213> Homo sapiens		
<220 ×		
<pre>&lt;223 &gt; Cdc25 hs ribozyme bind:</pre>	ing site	
±(400 × 3627		
aaattggata aaaatccaa	1	Ġ
±1210 > 3618		
8.1118 19		
S0128 DNA		
-12138 Home sapiens		
The Supering		
+1112008		
+0.23% Cdc25 hs ribozyme bind:	ing site	
1.75. Cdc25 is fibozyme bind.	ing site	
+:400:> 362:8		
	1	۲.
qataaaaatd caaacctag	1	9
15 5 5 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		
+010% 3629		
+:::11: 13		
HILLAN DNA		
<pre><pre></pre> <pre>Homo sapiens</pre></pre>		
·220		
<223> Cdd25 hs ribozyme bindi	ing site	
4		

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<pre>&lt;210 &gt; 3630 &lt;211 &gt; 19 &lt;212 &gt; DNA &lt;213 + Homo sapiens</pre>	
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<210 > 3631 <211 > 19 <212 > DNA <213 > Homo sapiens	
<220 > <223 > Cdc25 hs ribotyme binding site	
<400× 3631 gaagagattt cagatgaat	19
<pre>&lt;210 &gt; 3630 &lt;211 &gt; 19 &lt;210 &gt; PMA &lt;213 &gt; Homo sapiens</pre>	
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<pre>st220s st235 Cdc15 hs ribozyme binding site</pre>	
k4008 3833 dagatgaatt aatggagtt	19
+00.160+ 3.634 +00.110+ 14 +00.110+ DNA +00.130+ Homo sapiens	

<pre>+C220 * +C023 * Odc25 hs ribozyme binding site</pre>	
<pre>0400 + 3634 agatqaatta atggagttt</pre>	19
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<pre>&lt;220&gt; &lt;223&gt; Odc25 hs ribozyme binding site</pre>	
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+0210 + 3637 +0211 + 19 +0212 + DMA +0213 + Homo sapiens	
<pre><dd000 <d0005="" binding="" hs="" pre="" ribozyme="" site<=""></dd000></pre>	
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HILLON 3638 HILLON 19 HILLON DNA HILLON Homo sapiens	
H2230+ Cdc25 hs ribozyme binding site	
K400> 3638 tqqaqttttc cotqaaaqa	19

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<2105 3639
4211 - 13
<212 DMA
H213 + Homo sapiens
*(220 ×
H223 - Odc15 hs ribozyme binding site
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                                                                                             19
otgaaagato aagaagcaa
300109 3640
\pm 211 + 19
-1212 > DNA
+2213> Homo sapiens
<1220 >
%223> Odc25 hs ribozyme binding site
74005 3640
                                                                                             19
aaqtqgccta tatcgctcc
<.210 > 3641
<0.11 > 19
KO135 DNA
ACLIST Homo sapiens
4111 O +
+22.3 > Cdc25 hs ribozyme binding site
<400 > 3641
gtggddtata togdtoddd
                                                                                             19
HI210H 384I
4..11 • 19
40110 - INA
-1.13 Homo sapiens
\pm 2239 \cdot \text{Cdc.} 15 \text{ hs ribozyme binding site}
-1400b 3640
ggcctatate geteccegt
                                                                                             19
R01100 3643
+0.110 19
HI 110 DNA
\cdot \text{COLC} \cdot \text{Home sapiens}
<2210 ×

<
```

H4400 k 9643 tatatophto coogtogat	19
+1210 x 3:644 +1211 x 19 +1212 x DNA +1213 x Homo sapiens	
-120 ×	
+223 Cdc25 hs ribozyme binding site	
+:400 + 3644 geteccepte gatgocaga	19
-:2105 3645	
+:211> 19	
KODION DEL CONTROLLO DE LA CON	
+213> Homo sapiens	
+12.20>	
<pre><dd3> Odc35 hs ribozyme binding site</dd3></pre>	
-:400> 3645	
cagagaactt gaacaggcc	19
H210 × 3646	
<2111 × 19	
+1212% EMA	
+213 > Homo sapiens	
+1/20>	
MARIN Cac25 hs ribozyme binding site	
5(4)005 3646	
tqqaaaaatt caaggacaa	19
K0108 3647	
H11115-19	
HINLING LIMA	
+00130-Homo sapiens	
-ccos	
+2003+ Cacl5 hs ribozyme binding site	
-(400):- 3e47	
नवण्या वर्षात्रः वृष्वबद्धववःto aaggacaac	19
manua . 15 aaggacaac	13
-:::10:- 3648	
(2118-19)	
+:21/2: DNA	

<213> Homo sapiens	
-: <u>:::::::::::::::::::::::::::::::::::</u>	
40003 + Odd05 hs ribozyme binding site	
+:400 + 3648	
daadacaata ccagataaa	19
<0.10 + 3643	
-(211 + 19)	
HO10 + DNA	
+:213 - Homb sapiens	
<0.00 ·	
<pre><pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><td></td></pre></pre></pre>	
<400 · 3649	
ataccagata aagttaaaa	19
+(210 + 3650	
<211 + 19	
H212 + DNA	
x213 · Homo sapiens	
•	
<220 s	
<223 - Odc25 hs ribozyme binding site	
-(400)+ 3650	
ayataaagtt aaaaaaaag	19
<210% 3651	
<2215/19	
KORON DNA	
ALLINE Homo sapiens	
+(220)+	
<pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre>Cdc15 hs ribozyme binding site</pre>	
,	
<40.00 3651	
qataaagtta aaaaaaagt	. 19
HILLON 3651	
0.11:- 19	
HI 1.0 - DNA	
-Clist Bomo sapiens	
(C2.0)	
<22.00 Cdd25 hs ribozyme binding site	
<400> 3652	
aaaaaaagta tttttctgg	19

<pre>&lt;:210 + 3653 &lt;:211 + 19 &lt;:212 + DNA</pre>	
Home sapiens	
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-:400 - 3653 aaaaagtatt tittotggod	19
+C10 + 3654 +C211 + 19 +C212 + ENA +C213 > Homo sapiens	
<pre>&lt;220 / &lt;223 · Odc15 hs ribozyme binding site</pre>	
-:400 × 3654 aaaaatattt ttotggoda	19
+210% 3655 #211% 19 +211% DNA +211% Home sapiens	
-0000- -0003- Cdc05 hs riboxyme binding site	
<400> 3655 aaagtatttt totggooda	19
+1210+ 3656 +1211+ 19 +1210+ DNA +1213> Homo sapiens	
+22005 +2230 Cac25 hs ribozyme binding site	
+:4005 3656 aagtattiti otggodaag	19
+0110: 3+57 +011: 14 +010: DNA +013: Homo sapiens	
+:220 +	

-MARRY - Cdo25 hs ribozyme binding site	
त्र4 (म.) 3657 agrattitti tiggodaagg	19
<pre><pre><pre><cold +="" 365%<="" pre=""> <pre><cold +="" 19<="" pre=""> <pre><cold +="" dna<="" pre=""> <pre><cold +="" home="" pre="" sapiens<=""></cold></pre></cold></pre></cold></pre></cold></pre></pre></pre>	
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- (400 + 3658 augaaagoto aggaagggo	19
<pre>&lt;310 + 3659 &lt;311 * 19 &lt;312 * DNA &lt;313 * Homo sapiens</pre>	
<pre>k220 + k223 + Odd25 hs ribozyme binding site</pre>	
्र400% 3659 qqaaqgqqtt atgtttaaa	19
<pre>#210 - 3660 +2211 - 19 +2212 - DNA +2213 - Home sapiens</pre>	
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+02100 3661 +00100 19 +00100 DNA +00130 Home sapiens	
+02000 +02000 hs ribozyme binding site	
-:400% 3661 agcttatatt taaagaaga	19
+03100+ 3862 +03119-13	

HO12 + DNA HOMO sapiens	
+3200+ +323+ Odo25 hs ribozyme binding site	
-(400 - 0662 gottatgttt aaagaagad	19
KU10 + 3663 KU11 + 19 KU1U + DNA	
<pre> K223* Cdc25 hs ribozyme binding site </pre>	
<400> 3663 cttatgttta aagaagaca	19
<pre></pre>	13
<pre>%2308 %2308 %2308 Odc25 hs ribozyme binding site</pre>	
-4005 3664 маамасаять tetetgtgt	19
<pre><cli><pre><cli><pre><cli><pre><cli><pre><cli><pre><cli><pre><cli><pre><cli><pre><cli><pre><pre><cli><pre><pre><cli><pre><pre><pre><pre></pre></pre></pre></pre></cli></pre></pre></cli></pre></pre></cli></pre></cli></pre></cli></pre></cli></pre></cli></pre></cli></pre></cli></pre></cli></pre></cli></pre>	
+22008 +2230+ Cdc.15 hs ribozyme binding site	
+400:- 3e69 agadagtoto totgtgtga	19
<pre>%0.1008 3666 %0.1108 19 %0.1108 DNA %0.1308 Home sapiens</pre>	
+0210+ +0220+ Cdc15 hs ribozyme binding site	
+:400× 3666	

.

abagtotete tgtgtgaba	19
+0010 × 3667 +0010 × 10 +0010 × DNA	
±113 → Homo sapiens	
-0220 > 04025 hs ribozyme binding site	
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+010 + 3668 <011 > 19 <010 > DNA	
<213 Homo sapiens	
<pre>&lt;020&gt; &lt;020&gt; &lt;023   Odc25 hs ribozyme binding site</pre>	
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<010 + 3669 <011 > 19	
<pre>ROLL ONA ROMO sapiens</pre>	
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+D10+ 3670 +D11+ 19 +D1C+ DNA	
+C.130 Home sapiens	
-mag- -mag- Cdc25 hs ribozyme binding site	
-:400:- 3670 cattactato actoagatg	19
	17
+02100+ 3671 +0.110+ 19	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
•	

<220.	
+:223 + Odc25 hs ribozyme binding site	
+14+00 + 3671	
antatoanto agatgotgg	1 9
+210 + 3672	
H211 + 19	
ANC 11 % DNA	
+1115 · Homo sapiens	
·220>	
+223 + 0dc25 hs ribozyme binding site	
<400 + 3672	
gaggaagatt ctaaccagg	19
K210 x 3673	
<111 - 19	
H212 → DNA	
<pre>K213   Homo sapiens</pre>	
+12.10 +	
<pre><!--223 * Odo25 hs ribozyme binding site</pre--></pre>	
AND STORY	
+:400 + 3673	19
aggaagatto taaccaggg	1.7
+C10 ≠ 3674	
-:::::::::::::::::::::::::::::::::::::	
KINI IN ENA	
<pre>&lt;113&gt; Homo sapiens</pre>	
and nome suppose	
x(220)s	
+223: Odo25 hs ribozyme binding site	
<4000-3674	
gaagattota accaggggc	19
+D100+ 3675	
H211:- 19	
ROLD DNA	
RD130 Homo sapiens	
-22%.	
-2130 Cacl5 hs ribozyme binding site	
+:400::-3470	
qdaddtgat* ggtgatttt	19
was magazing garagazine	± = '
K210× 3676	

+331 + 19 +3212 + DNA +3213 + Homo sapiens	
R220 - R225 - Odc05 hs ribozyme binding site	
-:400 - 3676 attagtgatt tittocaagg	19
+0210+ 3677 <0211+ 19 +0213+ DNA +0213+ Homo sapiens	
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<pre>&lt;4005 3678 tqqtqatttt tocaaggta</pre>	19
<pre>\$0.10 × 3679 \$0.11 &gt; 19 \$0.11 &gt; DNA \$0.213 &gt; Homo sapiens</pre>	
-00205 -00205 Cacl5 hs ribozyme binding site	
k4005 3474 gqtqattftf ccaaggtat	19
*:210> 3480 *:211> 14 *:212> DNA *:213> Homo sapiens	
<pre>+IIII0:- +IX23&gt; Cdc25 hs ribozyme binding site</pre>	

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+11.00% +11.03 + Cdc25 hs ribozyme binding site	
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<pre>#210 + 3682 #211 * 19 #212 * DNA #213 * Homo sapiens</pre>	
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<pre>%210 * 3683 %211 * 19 %212 * ENA %213 * Homo sapiens</pre>	
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n400% 3683 babbaagath tgaagtatg	19
kD10 + 3684 kD11 = 19 kD10 = DNA kD13 + Himo sapiens	
-0.120 - -0.123 - Cdcl5 hs ribozyme binding site	
d4000-3684 atotqaaqta tgtcaaccc	19
0110 + 3685 0111	

<pre>Hill: Hill: H</pre>	
(400 - 3685 qaagtatqto aacccagaa	19
<pre>&lt;0.10   3686 &lt;0.11   19 &lt;0.212   DNA &lt;0.213   Homo sapiens</pre>	
<pre>k210 &lt;1113. Odc25 hs ribozyme binding site</pre>	
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R400 x 3687 ggotgootta otgtogggg	19
<pre>K210 + 3688 K211 &gt; 19 K212 &gt; DNA &lt;213 &gt; Homo sapiens</pre>	
<pre><i220 <="" pre=""> <pre></pre> <pre></pre> <pre>Cdc25 hs ribozyme binding site</pre></i220></pre>	
r4000-3688 cottactqtc ggggaagtt	19
+02100+ 3689 <02110+ 19 <02120+ DNA <02130+ Homo sapiens	
+CLIO:- +C2I3: Cdc25 hs ribozyme binding site	
-:400:- 368.4 cggggaagtt ccagggtot	19

<pre>#C210 + 3690 #C211 + 19 #C212 + DNA #C213 + Homo sapiens</pre>	
<pre>k220. k223. Odo05 hs ribozyme binding site</pre>	
(400 - 3690 gyggaagtto cagggtotg	19
#210 + 3691 #211 + 19 #212 + DNA #213 + Home sapiens	
<pre>&lt;000 + &lt;0003 + Odc05 hs ribozyme binding site</pre>	
k400+ 3691 ttobagggto tgattgaga	19
+210 + 3692 +211 + 19 +212 > 5NA +213 > Homo sapiens	
-220- -223- Cdc25 hs ribozyme binding site	
-0400 - 3892 gggtotgatt gagaagttt	19
H210 + 3693 H211 + 19 H212 + DNA H313 + Homo sapiens	
<pre><idd0+ <id0+="" binding="" cdc25="" hs="" pre="" ribozyme="" site<=""></idd0+></pre>	
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02100-3094 02110-19 02120-DNA 02130-Homo sapiens	
e2230 Cdc25 hs ribozyme binding site	

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tgagaagttt tatgtdatt	19
±210 + 3695	
<211 • 19	
HOLLY DNA	
<pre><pre><ii13 homo="" pre="" sapiens<="" •=""></ii13></pre></pre>	
•	
-(22a)	
<pre>&lt;:223 + Odc15 hs ribozyme binding site</pre>	
•	
-400 - 3695	
gagaagtitt atgtcattg	19
<210 > 3696	
4.311 × 19	
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KI13> Homo sapiens	
k020 ×	
<223 > Cac25 hs ribozyme binding site	
<400+ 3696	
agaagtttta tgtcattga	19
-210 × 3697	
+211 + 19	
0212 × DNA	
-213 Homo sapiens	
<.120 %	
<pre></pre> <pre></pre> <pre>Cdc.15 hs ribozyme binding site</pre>	
- 400 + 369°	
gittitatgite attgattgt	19
HEEDON 3698	
WILLY 19	
HILLION DNA	
HP150 Homo sapiens	
et Ann	
KAAOON	
CDDS CdcD5 hs ribozyme binding site	
(400) 3698 **********************************	19
tratqtcatr gattgtogo	13
-0.100-3694	
43.100 1992 43.110 19	
<pre><l12> DNA</l12></pre>	
ALICA DIVIN	

<pre>::213 · Homo sapiens</pre>	
-1125 -	
-023 - Cdc.5 hs ribozyme binding site	
-:400 + 3699	
ateattgatt gtogotato	19
CL10 + 3700	
√211 + 19	
KOLAN DNA	
<pre>+D13 + Homo sapiens</pre>	
K220 H	
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attyattyte getateeat	19
<310> 3701	
3.11> 19	
KD12> DNA	
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F2008	
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attqtoqbta tocatatga	19
H210H 3702	
S211 × 19	
KO125 DNA	
CD13 · Homo sapiens	
::12ō ·	
02230 Odc25 hs ribozyme binding site	
<400 · 3702	
tytogotato catatgagt	19
C210:-370?	
0.1118-19	
COLOR DNA	
C2130 Homo sapiens	
022009	
02230 Cdcl5 hs ribozyme binding site	
k400% 31103	
gotatocata tgagtatot	19

#2310 + 3704 #2311 + 19 #2313 + DNA #2313 + Home sapiens	
<pre>HARD + HARD + CdcA5 hs ribozyme binding site</pre>	
0400) - 3704 patatgagta totgggagg	19
<pre>4D10 + 3705 4D11 + 19 4D12 + DNA 4D13 + Homo sapiens</pre>	
<pre>&lt;220&gt; &lt;223 &gt; Odc15 hs ribozyme binding site</pre>	
<pre>&lt;400&gt; 3705 tatgagtate tgggaggae</pre>	19
+0010 > 3706 +0010 > 19 +0010 > DNA +0013 > Homo sapiens	
<pre>&lt;:220&gt; &lt;:223 + Odc25 hs ribbzyme binding site</pre>	
<400% 370% aggababato bagggagbb	19
+2108 3707 +2119 19 +2129 DMA +2130 Homo sapiens	
+00000 +00000 Cdc25 hs ribozyme binding site	
+14008 3707 agggagoott aaacttata	19
+02100+ 3008 +02100+ 14 +02100+ DNA +02130+ Homo sapiens	
·(220>	

<pre><d::23 binding="" hs="" odc25="" pre="" ribozyme="" site<="" ·=""></d::23></pre>	
स्व्यापः 3703 सम्बद्धाः accttatat	19
+:210+ 3709 +:211+ 19	
H212 + DNA	
+213 - Homo sapiens	
x(120 x	
-113 - Odc15 hs ribozyme binding site	
-:400 × 3709	
oottaaantt atatagtoa	19
<210 + 3710	
<211> 19 <210> DNA	
NDID K BOA K213> Homo sapiens	
Daptons	
F22009	
+333+ Odc25 hs ribozyme binding site	
+4005 3710	
ettäääetta tatagteag	19
<210 ≥ 3711	
<2115-19	
HEIDH DNA	
<pre>SU13: Homo sapiens</pre>	
<2208	
<pre>&lt;00035 Cdc05 hs ribozyme binding site</pre>	
<4400% 3713	
taaacttata tagtoagga	19
K210% 3712	
H2110-19	
HOLLOW DNA	
HO13: Homo sapiens	
-m20:-	
COLOR Cdc15 hs ribozyme binding site	
<pre></pre>	
aacttatata gtcaggaag	19
KD10:- 3713	
(211) 19	

-:212'+ DNA	
<pre><!--213 Homo sapiens</pre--></pre>	
•	
-:220 ·	
<pre>0.223    Cdc.15 hs ribozyme binding site</pre>	
Col. caons ne lizzenjmo kilmalng tito	
<400 · 3713	
ttatatagto aggaagaac	19
- I a ca cage a agga aga a caga a	
<(21) + 3714	
<111 - 19	
HOLD FONA	
4213 - Homo sapiens	
The fibility sapiens	
<230 /	
<pre><anual co<="" control="" td=""><td></td></anual></pre>	
ksv edebs hs tibozyme binding site	
<400÷ 3714	
	19
asgasctgtt taacttott	13
<pre>&lt;0.10 × 3715</pre>	
<011 × 19	
KORIO MARIA	
<pre><di13> Homo sapiens</di13></pre>	
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agaactgitt aacttottt	19
.54.0564.5	
+:210+ 3716	
<211 \ 19	
KOLLON DNA	
<pre>%113 Homo sapiens</pre>	
<220.	
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834000 - 3716	٦.
daactgttta acttettte	19
<pre><pre><pre><pre></pre></pre></pre></pre>	
NUT1:- 19	
CD1. IN ENA	
HIMISH Homo sapiens	
************************************	
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·ACO: 3717	

tgtttaactt ctttctgaa	19
<pre>%210 * 3718 %211 * 19 %212 * DNA %213 * Homo sapiens</pre>	
+320+ +323+ Odel5 hs ribozyme binding site	
+401 + 3718 quittaaptib titotgaag	19
+010 + 3719 +011 + 19 +010 + DNA +013 + Homo sapiens	
<pre><pre><pre><pre><pre></pre></pre><pre><pre></pre><pre></pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><!--</td--><td></td></pre></pre></pre></pre>	
<pre></pre> <pre>&lt;400+ 3719 ttaacttott totgaagaa</pre>	19
K210× 3720 K211× 19 K212× DNA K213× Home sapiens	
<pre>&lt;22008 &lt;22038 Cdd25 hs ribozyme binding site</pre>	
k4005 3720 taacttotti otgaagaag	19
+02100+ 3721 +02110+ 13 +02120+ DNA +02130+ Homo sapiens	
-0.000:0.000:0.000:-	
H4005 3721 aacttetite tgaagaage	19
+001(3 372. +001(3 19 +001(3 BVA) +000(3 saniens	

+:220 + -:233 + 2dc15 hs ribozyme binding site	
(400 + 3722) gaaquocato gtocottig	19
+:210% 37:03 +:211% 19 +:710% DMA	
<pre><li>113 Homo sapiens</li></pre>	
<pre>&lt;120 * &lt;223 * 0dc25 hs ribozyme binding site</pre>	
-400> 3703 goodateqte cotttggac	19
<pre>&lt;210&gt; 3704 &lt;211&gt; 19 &lt;212&gt; DNA</pre>	
+213> Homo sapiens	
<pre>&lt;:200* &lt;223* Odo25 hs ribozyme binding site</pre>	
4400% 3724 Atoqtobott tggacacco	19
+010 × 3725 ×011 × 19 ×010 × DNA	
S2135 Homo sapiens	
-0200	
64008 3725 togtocottt ggacacca	19
H2105 3706 H2115 19 H2105 DNA	
#2130 Homo sapiens	
HUZ 604	
<pre></pre>	
r40no 3726 qaaqagaata atcatogtg	19
201m. 3707	* >

<211 • 19	
HIB - DNA	
+0:13 + Homo sapiens	
-1.1.2 0 +	
-0.023 - 0.1525 hs ribozyme binding site	
+(400 + 3727	
qaqaataaso atogtgtto	19
, a, a a a a a a a a a a a a a a a a a	
±0.10 ± 3728	
+C11> 19	
SOLIC - DNA	
+213/ Homo sapiens	
1211 Hillio Supreme	
<220>	
<pre></pre> <pre>&lt;</pre>	
tels table no likelyme binding bitte	
<400> 3728	
aataatbato gtgttocac	19
and a second sec	
K210 S 3709	
<211 · 19	
S212S DNA	
k013% Homo sapiens	
The History Suprems	
K220 +	
<223 - Odc.15 hs ribozyme binding site	
22 Sas. o no libonyme binding ofte	
<4008-3709	
toatogtgtt. ccactgtga	19
· · · · · · · · · · · · · · · · · · ·	
<010 > 3730	
<.115 19	
KO108 DNA	
<pre>%213 * Homo sapiens</pre>	
nate in the captions	
+020 +	
-123 · Odc25 hs ribozyme binding site	
the state of the s	
+:400.+ 3730	
datoqtgtfo cactgtgaa	19
-00100+ 3731	
-0.110-19	
HUIII DNA	
:::12	
1.2. Home Suprons	
+:22,00+	
and such in the short of the structure of the	

<pre>(400) 1731 astigtgaatt ctcctcaga</pre>	19
0.010 + 2732 0.011 + 19 0.012 + 1NA 0.013 + Homo sapiens	
<pre>&lt;:220 - &lt;:223 - Cdc25 hs ribozyme binding site</pre>	
0400 + 2732 otqtqaatto tootoagag	19
#210 + 3733 #211 + 19 #212 + DNA #213 + Homo sapiens	
<pre>&lt;210 - &lt;223 - Cdc25 hs ribozyme binding site</pre>	
<pre>-:400 + 3733 gtgaattoto otoagagāg</pre>	19
H210 + 3734 +211 + 19 +212 + DNA +213 + Hom: sapiens	
<pre>+:220* +:223* Cdc25 hs ribozyme binding site</pre>	
0.400 - 3734 aattotooto agagaggg	19
<pre>8210 + 3/35 +M11 + 19 8M11 + DMA +M11 + Home sapiens</pre>	
-uhnu- -uhnu- Odc25 hs ribozyme binding site	
+:400:+ 3735 taccactite tgcgtgaag	19
+0.10+3038 +0.10+1* +0.210+EMA +0.215> Humo sapiens	

$\pm 0.20$ . $\pm 0.23$ hs ribozyme binding site	
्याण + 3736 चापुबलबुक्टा totgaacca	19
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
+C1138 DNA +C1139 Homo sapiens	
<pre>&lt;:220* &lt;123* Odo25 hs ribozyme binding site</pre>	
<pre>&lt;400&gt; 3737 gacaggtoto tgaaccagt</pre>	19
<pre>&lt;:210&gt; 3738</pre>	
<2211> 19	
HIGHES AND KINDS	
<pre>&lt;213&gt; Homo sapiens</pre>	
<220>	
<pre>&lt;223&gt; Odc15 hs ribozyme binding site</pre>	
+:400> 3738	
tgaaccagta tootgoatt	19
- 11 us - 17 10	
<pre>&lt;210&gt; 3739 </pre>	
+2012 DNA	
-1313> Eomo sapiens	
<pre>02235 Cdc15 hs ribozyme binding site</pre>	
+:4008-3739	
aaccagtato ctgcattgt	19
+:210> 3740	
82211> 19	
4.2125 ENA	
::11:5 Homo sapiens	
-compas	
Hall ab Cdc.5 hs ribozyme binding site	
<:400> 3740	
atcotgoatt gtactacco	19

<pre>#210 &gt; 3741 #211 + 19 #212 + DMA #213 + Homo sapiens</pre>	
+1320 + +1323 + Cdc.5 hs ribozyme binding site	
-0400 - 3741 otgoattgta otacocaga	19
<pre>0016 - 3740 -011 - 19 -011 - DNA -013 - Home sapiens</pre>	
<pre>&lt;0.00 + &lt;0.03 &gt; Cdc.05 hs ribozyme binding site</pre>	
<400 - 3740 cattgtacta cocagaget	19
<pre>&lt;210 - 3743 &lt;211 - 19 &lt;213 - Bomo sapiens</pre>	
<2205 <223 Cdc25 hs ribozyme binding site	
<400 · 3743 decayageta tatateett	19
<pre>KD10 + 3744 KD11 &gt; 10 KD12 + DNA KD13 + Homo sapiens</pre>	
<pre>&lt;:220 - &lt;:223 - C4c15 hs ribozyme binding site</pre>	
<pre>&lt;4000 - 3044 dagagetata tateettaa</pre>	19
+U 1()+ 3 (4) +U 110-19 +U 110- DMA +U 150- Homo sapiens	
+0.20+ +0.25+ Cdc25 hs ribozyme binding site	

-:400 - 3745 gagstatata teettaaag	19
+210 + 3746 +211 + 19 +212 + DNA	
+213 · Homo sapiens	
<pre><dable color="block"><dable color="block"></dable><dable color="block"></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable></dable><td></td></pre>	
<pre>&lt;400 + 3746 qctatatata cttaaaggc</pre>	19
+210+ 3747 <211+ 19	
<212> DNA	
+:213> Homo sapiens	
<223 > Cdc35 hs ribozyme binding site	
×400 × 3747	
atatateett aaaggegge	19
+:210 + 3748 +:211 + 19	
<pre>&lt;212 + DNA</pre>	
<pre>4213 * Homo sapiens</pre>	
<220 %	
<pre>&lt;223 * Odc25 hs ribozyme binding site</pre>	
+(400 + 3748	
tatatoetta aaggogget	19
HZ10 + 3749	
*Ells 19	
+:21:0+ FNA +:21:0+ Homo sapiens	
k(220)-	
+CCCC CdcC5 hs ribozyme binding site	
H40GH 3T49	
äagqdggdta cagagactt	19
H:100-3150	
+17.11(+ 1.9)	
K2120 DNA	

::213 > Homo sapiens	
+1220 ·	
H1223 - Cdc25 hs ribozyme binding site	
+(400 + 3750	
abagagaett ettteeaga	19
-U 10 + 3751	
+1.11 + 19	
HILLS DNA	
HO13 + Homo sapiens	
+:220 ·	
+:223 > Cdc25 hs ribozyme binding site	
<400≻ 3751	
cagagactto tttocagaa	19
<210> 3752	
<211 > 19	
<pre>&lt;212   DNA &lt;213   Homo sapiens</pre>	
C213 / Rollio Sapiens	
k:220>	
<pre>&lt;223&gt; @dc25 hs ribozyme binding site</pre>	
Alexander Communication Commun	
<pre>4400 &gt; 3751 gagacttett tocagaata</pre>	19
Mayasetset teeagaata	10
+:::10 × 3753	
40115-19	
+:212> DNA	
+2135 Homo sapiens	
+:22.0 ×	
-0023 - Cac25 hs ribozyme binding site	
<:400% 375%	
agact.tottt ocagaatat	19
H210H 3054	
K2115-19	
HC12N ENA	
+2013> Homo sapiens	
-(MMO)-	
+2003+ Cdc25 hs ribozyme binding site	
4400≥ 3754	
gacttctttc cagaatata	19

+010 + 3755 +011 + 19 +011 + DNA +013 + Homo sapiens	
+3220 + +3223 + Odc.5 hs ribozyme binding site	
0400% 375% ttppagaata tatggaact	19
+0010 + 3756 +0010 > 19 +0010 > ENA +0013 > Homo sapiens	
<pre>&lt;2206&gt; &lt;223&gt; Cdc25 hs ribozyme binding site</pre>	
-:400> 3756 ccagaatata tggaactgt	19
<pre>&lt;210 &gt; 3757 &lt;211 &gt; 19 &lt;212 &gt; DNA &lt;212 &gt; Homo sapiens</pre>	
<pre>&lt;:220&gt; &lt;:223&gt; Cdc25 hs ribozyme binding site</pre>	
<400> 3757 cacagageta etgecetat	19
<pre>%210 * 3/54 %211 * 19 %212 * DMA %215 * Homo sapiens</pre>	
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>v4008 3758 tactdochta tgcatcatc</pre>	19
+021(0+ 3/54 +0210+ 14 +0010+ DMA +00130+ Home sapiens	
+0220×	

$\cdot:$ 223 $\cdot$ 3dc35 hs ribbzyme binding	site
±1400 + 3759	
octatgoato atcaggacc	19
<210 + 3760	
4211 + 13	
H212 + DNA	
+C213 - Homo sapiens	
*1220 * *22.2 ** ** ** ** ** ** ** ** ** ** ** ** **	
<pre>&lt;2233 + 0de25 hs ribozyme binding</pre>	site
<400 + 3160	
atgeateate aggaecaea	19
<210 + 3761	
<211 ÷ 19	
K212 - DNA	
<213> Homo sapiens	
.55	
<220 × <223 · Odc25 hs ribezyme binding	aito
Kills Codols has riberyme binding	Site
<400> 3761	
agactgagtt gotgaggtg	19
<210 × 3762	
<211.5 19	
<21.05 DNA	
<2138 Homo sapiens	
k(226)5	
<pre>&lt;2230- Odc25 hs ribczyme binding</pre>	site
k(400)S   3762	
cidadgidic gaagccaga	19
+(2.10) + 3065	
H211H 19 H212H DNA	
KIII ONA KIII Homo sapiens	
allo memo sapiens	
+:200:+	
<pre>02130 Cdc25 hs riboxyme binding</pre>	site
-:4000 - 30 <b>63</b>	
qqaqqaqatt goodttotg	19
-0.110031164	
K2112 19	

RC103 DNA	
H213 · Homo sapiens	
(220) •	
HAAR - OdoN5 hs ribozyme binding site	
and sales is timely will be seen as a second	
₹400 + 3764	
	19
gattqccst.ctggtgaag	1 2
<010 + 3765	
<pre><pre><pre></pre></pre></pre>	
HILL HILL HILL HILL HILL HILL HILL HILL	
化1.13 · Homo sapiens	
<2.10 %	
<pre><dul><!-- Cdc. S hs ribozyme binding site</pre--></dul></pre>	
<400 · 3765	
attqccttc tggtgaagg	19
<210 > 3766	
<211≥ 19	
KOIDE DNA	
<213 - Homo sapiens	
ALTO HOME Suprone	
<000%	
<2003 - Ode05 hs ribezyme binding site	
Seller Casho its libertyme binding site	
<400 · 3766	
goodatgata acattodag	19
ghobacyaca acaccecay	1
<210 · 3767	
<2115-19	
SOLD - DNA	
40.13 · Homo sapiens	
<220 ·	
<pre>&lt;123% Ode.5 hs ribezyme binding site</pre>	
<4000 3767	
tgafaacatt, ccagccact	19
<2100 3768	
KD119-19	
KULTUR DNA	
-CCLO- Homo sapiens	
•	
$\star(Z_{+}^{*},0)\star$	
<pre></pre> <pre></pre> <pre></pre> <pre></pre> <pre>Cdc25 hs ribozyme binding site</pre>	
2. Sando no razoujmo ornarny oros	
< 4000 3768	

gataacatto cagcoactg	19
+:210 + 3769	
-0.211 • 19	
<pre>&lt;011 + DMA</pre>	
<pre><d:13 -="" homo="" pre="" sapiens<=""></d:13></pre>	
+220 +	
<pre></pre>	
<400 + 3769	
onggotgots acaagtoac	19
<2.10 + 3.770	
K221 × 19	
HILL FORA	
<pre>&lt;0.13&gt; Homo sapiens</pre>	
+:220 ×	
+223> Odou5 hs ribozyme binding site	
<400 ≥ 3770	
otascasqtc accaaaaag	19
k010+ 3771	
<211> 19	
RDID + DNA	
s213 m Home sapiens	
+220 +	
<pre>&lt;2003 Odc05 hs ribozyme binding site</pre>	
-:400s 3771	
aagadgsett etggatgge	19
HIL10H 377/	
H211N 19	
ROMA DNA	
RM130 Homo sapiens	
-00.008	
CMC130 Cdc15 hs ribozyme binding site	
-:400:- 3772	
acccaagatt attaaaaga	19
(C10)- 3773	
40117-1+	
HILLING DNA	
<2213> Homo sapiens	

HARRO - HARROS HS ribozyme binding site	
-(4)0 - 3773 cocaagatta ttaaaagat	19
+1210 + 3774 +1711 + 19	
HIMI - DNA HIMI - Homo sapiens	
+:220 +:223 + 04c25 hs ribozyme binding site	
+400> 3774 caagattatt aaaagatgt	19
<pre>&lt;210&gt; 3775 &lt;211&gt; 19 &lt;212&gt; DNA &lt;212- Constant</pre>	
<pre>&lt;213 &gt; Homo sapiens &lt;220 &gt; &lt;223 &gt; Cdc25 hs ribozyme binding site</pre>	
<pre>&lt;400 &gt; 3775 aagattatta aaagatgto</pre>	19
<pre><i210 -="" 3776="" <i211=""> 19 <i212> DNA </i212></i210></pre> <pre>&lt;213 &gt; Homo sapiens</pre>	
<pre>%AndOpe %AndOpe %</pre>	
+400× 3°7% aaaaqatgto totgoaaao	19
+D100+ 3777 +D11+ 1+ +D110+ DNA	
<pre>SUL130 Homo sapiens <pre>su2000</pre></pre>	
$\pm 20.30^\circ$ Cdc.5 hs ribozyme binding site	
<pre>+(40.00 377)* aagatgtoto tgcaaacca</pre>	19
+:210> 3778	

-:211 + 19	
S213 + DUA	
-0.213 · Homo sapiens	
+:22♥+	
$< 223 \cdot \text{Cdc.}25 \text{ hs ribozyme binding s}$	ite
+(400 + 3778	
osacaggota ccaapttgt	19
H010 + 3779	
+:211 + 19	
HOLD - DNA	
HDMO sapiens	
<:200>	
<223> Odc.5 hs ribozyme binding s.	ite
-:400> 3779	- 0
ctaccaacti gtatccagg	19
<210> 3780	
<211> 19	
SCILS DNA	
<pre><d13> Homo sapiens</d13></pre>	
25565	
<220>	
<223> Odc15 hs ribozyme binding s.	ite
₹400 > 3780	
odaaottgta todaggoot	19
Solda Structure and Colorado C	19
s:::10 × 3781	
<0.11 × 19	
SC12 S DNA	
<pre>%313 * Homo sapiens</pre>	
The state of the s	
K12203	
<pre>%223 · Odc25 hs ribozyme binding s:</pre>	ite
+:400+ 3781	
aacttgtate caggeotgg	19
, , , , , , , , , , , , , , , , , , ,	
+M10> 3780	
-M11> 1+	
HOLD DNA	
-1 18 Homo sapiens	
•	
+17.20c+	
<pre>:223&gt; Cdc25 hs ribozyme binding s:</pre>	ite

ार्था । 3782 चुत्रुवक्षां ggatt aggittcag	19
H210+ 3783 H211+ 19 H212+ DNA	
<pre>Homo sapiens</pre>	
+220 + +223 + Odc25 hs ribozyme binding site	
-1400 - 3783 gaatggatta ggtttcagc	19
S210 + 3784 4211 × 19 4212 + DNA	
<213> Homo sapiens	
<pre>k220s k223 + Cdc25 hs ribozyme binding site</pre>	
<4000 3784 ggattaggtt teageagag	19
<pre>&lt;210&gt; 3785 </pre> <pre>&lt;211&gt; 19 <pre><pre><pre>&lt;210&gt; ENA</pre></pre></pre></pre>	
<pre>4213&gt; Homo sapiens</pre>	
<pre>&lt;2000 &gt; &lt;2003 &gt; Odc05 hs ribozyme binding site</pre>	
<400 > 378 %	10
qattaggttt cagcagagc	19
+0.10+.3786 +0.11+.19	
HILLS DNA	
<pre>Hills Himo sapiens</pre>	
+C220+ +C223+ Cdc. 5 hs ribozyme binding site	
+1400th 3784	1.0
attäggttto agcagaget	19
+0.160+ 3780 +0.110+ 13	
-0:12:- DNA	
<pre>+C2130 Homo sapiens</pre>	

<pre>&lt;:220&gt; <!--:223 + Odc25 hs ribozyme binding site</pre--></pre>	
H400 + 3787 tyggagagto otggagotg	19
HC210 + 3789 HC212 + DNA HC213 + Homo sapiens	
+:220 + +:223 + Odc25 hs riboxyme binding site	
<4400> 3788 gagetggete tataaggea	19
<pre>&lt;210 &gt; 3789 &lt;211 &gt; 19 &lt;212 &gt; DNA &lt;213 &gt; Home sapiens</pre>	
<pre>&lt;220 + &lt;223&gt; Odc25 hs ribozyme binding site</pre>	
k400 + 3789 gutagotuta taaggoago	19
<pre>KD10 + 3790 KD11 + 13 KD1D&gt; DNA KD13 + Homo sapiens</pre>	
+02200+ Cdcl5 hs ribozyme binding site	
-:400:- 3790 tqqqtotata aggcagoot	19
+02100+ 3791 +02115 14 +02125 DNA +02135 Homo sapiens	
<pre>&lt;:220:- &lt;:223: Cdc25 hs ribozyme binding site</pre>	
+:400:-3791 aggcagectt gagttgcat	19

-:210 ·	3792		
41.11 ·			
-1.111 ×			
	Homo sapiens		
1.1.	I. M. / Sapiens		
-1220 -			
	~	1.1	
	Cdc. 5 hs ribozyme bi	naing s	site
-(400 -			
qaqttq	gcata gagatttgt		19
·1110 ·			
<pre><!--!!!!</pre--></pre>	1일		
$\cdot \left[ \begin{array}{ccc} & & & \\ & & & \\ & & & \end{array} \right] \cdot \left[ \begin{array}{ccc} & & & \\ & & & \\ & & & \end{array} \right]$	Z(1127		
<1.13 +	Homo sapiens		
<2000 S			
<323 €	Cdc25 hs ribozyme bin	nding s	site
	1	,	
<400×	3793		
	agatt tgtattggt		19
	ighter egedeegge		
<:2105	379.1		
d2113			
:2115			
	Homo sapiens		
. 5. 5. 5			
H2200	21.55	, .	
41233	@dcl5 hs ribozyme bir	nding s	site
<[400]>			1.0
atagag	gattt gtattggtt		19
<0.105			
s:::117 >			
·2213 ·	Hime sapiens		
K2200.+			
-12233	@doff5 hs ribozyme bir	nding s	site
(400)	3795		
qagat t	itgta ttggttcag		19
<del>-</del>			
.:::1(::-	3.196		
-00125-			
	Homo sapiens		
	oup tono		
·00200			
	Cdc25 hs ribozyme bir	ndina (	site
~~~ 0/	careed to tinosyme bit	maring 3	

(4)0 + 3796 partogtato ggttcaggg	19
#210 # 3797 #211 * 19 #2213 * DNA #213 * Homo sapiens	
+320 + +323 + Odc25 hs ribozyme binding site	
k(400 + 3797) tgtattgqtt cagggaact	19
<210> 3798 <211> 19 <210> DNA <213> Homo sapiens	
<pre>&lt;2000 class</pre>	
<400% 3798 qtattggttc agggaactc	19
<pre>KD10: 3799 KD11: 19 HD12: DNA HD13: Homo sapiens</pre>	
<pre><d200% <="" <200%="" pre=""> <pre><dc25 binding="" hs="" pre="" ribozyme="" site<=""></dc25></pre></d200%></pre>	
04008 3799 dagggaacto tggcattos	19
+00100+ 3500 +00110+ 19 +00120+ DNA +00130+ Homo sapiens	
<pre><pre><i2200< pre=""> <pre><i2230 binding="" cdc25="" hs="" pre="" ribozyme="" site<=""></i2230></pre></i2200<></pre></pre>	
+:400:- 3M00 chotagoatt cottttoco	19
+00100+ 3801 +00110+ 104 +02120+ ENA	

+C213 + Homo sapiens	
(2.20)	
-0223 - Odc.25 hs ribozyme binding site	
+400 + 3801	
totypoatto ottitocca	19
+(210 + 3802	
±2.11 + 19	
HOLD - DNA	
HOME Sapiens	
+1220 +	
<pre></pre> <pre></pre> <pre>Cdc25 hs ribozyme binding site</pre>	
<400.0 3802	
ggcattoott ttoccaact	19
<210> 3803	
<211> 19	
+:01.3 > DNA	
<pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre></pre></pre>	
H2008	
+203 * Cdc25 hs ribozyme binding site	
·400 · 3803	
deatheath tecesacte	19
K210% 3804	
s211 + 19	
KC125 DNA	
$\pm 213 \pm  ext{Homo}$ sapiens	
√200 ÷	
+0003+ Odc05 hs ribozyme binding site	
44000-3804	
pattecttut decaacted	19
H210N 3805	
-D118 19	
KULI DE CNA	
-Clish Homo sapiens	
::D1 us-	
H2230+ Odc25 hs ribozyme binding site	
-:400% 3805	
atteetttte ceaacteet	19

<210> 3306	
H211 + 19	
RC12 - DMA	
<pre>&lt;:213 → Homo sapiens</pre>	
s(220 s	
<pre><!--223 + Cdc25 hs ribozyme binding site</pre--></pre>	
+:400.+ 3806	
ttoocaseto eteatgtet	19
-210× 3807	
+211> 19	
HD1D> ENA	
+213> Homo sapiens	
<:220>	
<pre><d223> Cdc25 hs ribbzyme binding site</d223></pre>	
-:400 ≥ 3807	
coascicctc atgictici	19
-1.10 × 3808	
<2118 19	
KO12N DNA	
+313 · Homo sapiens	
<22.0 s	
<pre><!--D3* Cde.5 hs ribozyme binding site</pre--></pre>	
k400% 3808	
tectoatgic ticteacaa	19
K210 % 3809	
K2115-19	
HILLIN DUA	
<pre><!--!!30 Homo sapiens</pre--></pre>	
(200) ·	
<pre>*223 * Cdcl5 hs ribozyme binding site</pre>	
-:400:- 3809	
stoangtht, ctcasaago	19
H210H 3810	
00118-19	
CUITED DNA	
:2213: Home sapiens	
.:25fr.	

-0.223 - Odc25 hs ribozyme binding site	
-:400 + 3910	
toatqtotto toacaagoo	19
H2109 3811	
(211)/19	
KALAS DNA	
213> Homo sapiens	
<pre>&lt;223 Cdc25 hs ribozyme binding site</pre>	
*225 * Odd25 Hs Tibblyme binding Side	
-400 × 3811	
atgtottoto acaagocag	19
√210≥ 3612	
<211× 19	
HIDE DNA	
<213 Momo sapiens	
213 mm oup 10mo	
<:2.10 ×	
<233 - Odc25 hs ribozyme binding site	
<400 → 3810	
dagodaacto tittototot	19
	1.2
€210 + 3813	
(211 × 19)	
KO12 HIDNA	
<pre>&lt;013 &gt; Homo sapiens</pre>	
<220.5	
<2003 Cdc.5 hs ribazyme binding site	
KL. 3 · Cdc. 5 hs fibelyme binding site	
<400 × 3813	
godakotott totototgg	19
·210:-3814	
- 2010 - 19	
CHILLY DNA	
CDLSC Homo sapiens	
- min co-	
HIM 30 CdcU5 hs ribozyme binding site	
4340CC 3814	
deaadte;tt dtetetggg	19
-:::10:- 3815	
H2111: 19	
New Artist Control of the Control of	

HORIZ + DNA HORO sapiens	
+:22% + Cdc25 hs ribozyme binding site	
-:4000-3815 	19
+1210 + 3816 +1211 + 19 +1212 + EMA	
<pre>St213 - Homo sapiens</pre>	
<223 / Cdc25 hs ribozyme binding site	
<400 > 3816 actotttoto totgggott	19
<pre>k010 + 3017 k011 + 19 k010 + DNA k013 + Homo sapiens</pre>	
\$0.00 ×	
FURBLE Cdc. 5 hs ribozyme binding site	
k400× 3817 totttatata tgggattag	19
<pre>&lt;210&gt; 3818 &lt;211&gt; 19 &lt;0012&gt; EMA</pre>	
+2135 Home sapiens	
<pre>+00.008 +00.038 Cdc.05 hs ribozyme binding site</pre>	
ki460i- 3818 ototoggott ogggotatg	1 <sup>ç</sup>
<pre>*D100 3819 +D110 19 *D100 DMA</pre>	
+2130 Homo sapiens	
<pre>+CMCO+ +CMCO+ Cdc(5 hs ribozyme binding site</pre>	
+:400% 381 <del>9</del>	

.

totgggotto gggotatgo	19
<:310 + 3820	
3.13 - 3.27 3.211 - 13	
HU12 - DNA	
-NN13 - Homo sapiens	
<.220 ≥	
<pre>0223 * Odc25 hs ribozyme binding site</pre>	
<400 - 3820	
ottogggota tgcaagago	19
KC10 + 38C1	
<211 + 19	
AGION ANG AGION	
<pre>&lt;213 Homo sapiens</pre>	
K220 +	
<2003 > Cdc25 hs ribozyme binding site	
₹400 / 3811	
paaqagogtt gtotacott	19
K010% 3802	
<2115-19	
SELLES DNA	
<pre><!-- Clide Homo sapiens</pre--></pre>	
+:220+	
<pre></pre> <pre>C223&gt; Ode25 hs ribozyme binding site</pre>	
<400> 3822	
gagegttgto taccttott	19
gagogetg. We cadoccook	1.
<210 + 3%23	
KIII + 19	
H212 + EMA	
R213: Hemo sapiens	
+:22ti:-	
RCLN: Cdc. 5 hs ribozyme binding site	
+:400:3603	
	19
quightightal contentito	17
HIL10H 3814	
30110-19	
HURLING DNA	
+2112 Homo saniens	

$<$ 220 $\cdot$ $<$ 2223 $\cdot$ Cdz25 hs ribozyme binding site	
·(40) + 3824	
tytotacott officity	19
<210 + 3825	
s(211 + 19)	
HO12 HONA	
<pre>-C13 + Homo sapiens</pre>	
+:220 +	
+0.03 + Odo25 hs ribozyme binding site	
-:400 × 3825	
gtotacette tttetttgt	19
<210 × 3826	
<211 × 19	
<pre>&lt;210&gt; DNA</pre>	
<pre>&lt;113 Homo sapiens</pre>	
K220%	
<pre>&lt;223 * Cde25 hs ribozyme binding site</pre>	
<400 × 3826	
ctaccttett tetttgtat	19
<210 · 3817	
K2115-19	
KBIBH DNA	
<2135 Homo sapiens	
<008	
-CDD3+ Cdc25 hs ribozyme binding site	
<4000 3807	
tagnttstit cittgtatt	19
H210M 3808	
(III 118-19	
HUILDN DNA	
-Wlib- Homo sapiens	
kii. ex	
-0.35 Odc25 hs ribozyme binding site	
-:400:- 3628	
anditotifo titgiatit	19
<210x 3829	

÷211 · 13	
<pre><!--212 + DNA</pre--></pre>	
<pre><pre></pre> <pre><pre></pre> <pre></pre> <pre>####################################</pre></pre></pre>	
A.T. HAMA Suprono	
×(120 ·	
-1.1.23 · Cdc.15 hs ribozyme binding site	
(100 - 100 )	
(4.00 ± 362.)	1.0
official to the total to	19
H210 × 3830	
+2211 + 19	
SU 10 + DNA	
+213 - Homo sapiens	
<22.00	
<pre>&lt;2.13 · Odc.15 hs ribozyme binding site</pre>	
•	
<400> 3830	
ttotttottt gtattttoo	19
<210× 3831	
<211> 13	
<212> DNA	
<213> Homo sapiens	
Collar Months Saprens	
2110.	
<220 \	
<223 · Odc15 hs ribozyme binding site	
₹400 ± 3831	
tttetttgta tttteette	19
K210 + 3832	
-0115 19	
HITTS DNA	
-1113 · Homo sapiens	
-02.2.0 h	
-C233 Odc25 hs ribozyme binding site	
K400 + 3932	
tettigtatt tiecticii	19
-07107-3833	
*C110: 13	
00100 DNA	
*ILIDA Homo sapiens	
+0000+	
<223> Cdc25 hs ribozyme binding site	

.

<pre>d400 + 3033 objects toottottt</pre>	19
H210 + 3834 H211 + 19	
+C212 + DNA +C213 + Homo sapiens	
+0000 + +0003 + Odc25 hs ribozyme binding site	
<pre>k400 - 3834 tttgtatitt cottotttg</pre>	19
H210 H 3835 K211 H 19	
<pre>&lt;213&gt; DMA &lt;213&gt; Homo sapiens</pre>	
<2208 <223 Cdc25 hs ribozyme binding site	
<pre>k400 * 3835 ttgtattite offcttgt</pre>	19
H210> 3836 H211: 19	
<pre>COID&gt; DNA COID&gt; Home sapiens</pre>	
<pre>+0200 &lt;223&gt; Cdcl5 hs riboxyme binding site</pre>	
H4000-3836 tattitiooti ottigitto	19
+01\day 3837 +01108 19	
+:Clib+ DNA +:Clib+ Hemo sapiens	
+:2200+ +:2230 Cdc.)5 hs ribozyme binding site	
k4000-3837 attitocito titgittoo	19
+0.100+ 383+ +0.110+ 14	
+C2130 DMA +C2130 Homo sapiens	

·(22.) ·	
+:223 + Cdc.25 hs ribozyme binding site	
-1400 # 3838	
tttoottott tgtttooco	19
<:210 + 3839	
-:211 - 19	
HOLD - DNA	
+1.13 + Homo sapiens	
+12.20 s	
+223 · Cic25 hs ribozyme binding site	
Selection of the Tibozyme binding site	
<400 > 3839	
tibottotti gittocccc	19
-2102242	
<2210 > 3840	
<pre>80119 19 &lt;010* DNA</pre>	
<213> Homo sapiens	
S2137 Hollio Sapiens	
<220 -	
<pre>&lt;2233 Cdc25 hs ribozyme binding site</pre>	
-1100 - 20A	
(400) 3840	19
attatttytt teceestet	19
k(210 × 3941	
+(211 + 19	
N212 DNA	
<213 · Homo sapiens	
K2.00 +	
<pre><pre></pre>% Cdc15 hs ribozyme binding site</pre>	
s(400 × 3841	
ttotttattt coccetett	19
80:10 + 3842	
HD11:- 19	
HILLION DNA	
+2.13 · Home sapiens	
+:imc:+	
+2030 Odc05 hs ribozyme binding site	
Table Guest No Tibozyme binding Sice	
H400V 3842	
tetttgttte cecetettt	19

HC210 + 3843 HC211 + 19 HC212 + FNA HC213 + Homo sapiens	
+1220+ +1227+ Cdc25 hs ribozyme binding site	
HQD() = 3843 httpsccstd titlettiti	19
+D10> 3844 +D11> 19 +D12> DNA +D13> Homo sapiens	
<pre><dalor< td=""><td></td></dalor<></pre>	
+400 > 3844 teaccotott tottttta	19
+1210 > 3845 +1211 > 19 +1012 > DNA +1213 > Homo sapiens	
<pre>+220* +223* Cdc35 hs ribozyme binding site</pre>	
-:400 / 3845 cocceptettt etttttaa	19
+22100-3846 +2211-19 +22125 DNA +22135 Homo sapiens	
-1100. -11100 Cdc15 hs ribozyme binding site	
-:400:- 3346 enentetite titittaaa	19
+110: 3847 +111: 19 +111: DNA +113: Homo sapiens	
(U20) (U23) Cdc25 hs ribozyme binding site	

-:400 · 3847	
pototttott ttttaaaaa	19
<210 ← 3848	
<211> 19 <212 - 202	
+1212 × DNA	
+2213 Homo sapiens	
+:220 S	
<pre>&lt;1223 &gt; Cdc25 hs ribozyme binding site</pre>	
·:400 > 3848	
	1.0
ototttottt tttaaaaat	19
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<211> 19	
<212> DNA	
<213> Homo sapiens	
<0.00 >	
<pre>&lt;3&gt; Cdc25 hs ribozyme binding site</pre>	
william odd25 H5 TIBODYMC BINGING SICC	
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totttotttt ttaaaaatg	19
<010> 3850	
4211> 19	
S212> DNA	
<pre>&lt;213&gt; Homo sapiens</pre>	
<2208	
<pre>&lt;223&gt; Cdc25 hs ribozyme binding site</pre>	
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otttottttt taaaaatgg	19
K2108 3851	
<211> 19	
HILL DIA	
-213 · Home sapiens	
<220>	
CD238 Cdc25 hs ribozyme binding site	
(400): 3551	
titottitt aaaaatgga	19
The model of adducting a second of the secon	¥ J
₹£100-3452	
K211:- 19	
<2120 DNA	

HOMO sapiens	
<pre>&lt;2.00 - &lt;2.23 - Cic.05 hs ribozyme binding site</pre>	
-:400 - 3852 ttottttta aaaatggaa	19
+010 + 3853 +011 + 19 +010 + DNA	
<pre>%213 - Homo sapiens</pre>	
+3220 + 4223 + Odc25 hs ribozyme binding site	
k(400 - 3853 tggaaaaata aacactaca	19
<210> 3854 <211> 19	
<pre>&lt;012&gt; DNA &lt;013&gt; Homo sapiens</pre>	
Nation Remonsaprens	
<2230> <223> Cdc.:5 hs ribozyme binding site	
+:400 × 3854	
ataaacasta cagaatgag	19
<pre>&lt;:210 &gt; 3855</pre>	
<pre>%211&gt; 19 %212&gt; DNA</pre>	
<213 > Homo sapiens	
H220 h	
<pre>SDD3 - FCNA HH ribozyme binding site</pre>	
+:400 × 3955	
ttaaacgatt gcaggcgta	19
XC10:- 3656	
<2110-19	
<pre>&lt;012&gt; DMA </pre>	
-:22().	
+:22N+ PCNA HH ribozyme binding site	
+:400:- 3856	1 ^
tgcaqgcqta gagagtggt	19

R210+ 3857 R211+ 13 R212+ DNA			
+:213 + Homo	sapiens		
KU20 +			
+1223 + F*TNA	HH ribozyme	binding	site
·:400 · 3857			
qaqaqtqqtc	gttgtcttt		
-1110 - 3858			
4211: 19 4212: DNA			
<212 - D.A <213 - Homo	sapiens		
<220>			
	HH ribozyme	binding	site
<:4005 3858			
aqtqqtcqtt	gtettteta		
<210≻ 3859			
K211 - 19			
SIDIDE DNA			
<1.13 m Homo	sapiens		
· [][[][[]			
HIII3 - FCNA	HH ribozyme	binding	site
<400> 3859			
gatoattato	tttctaggt		
-121d> 3860			
02115 19 02175 ENA			
-IT13> Eomo	sapien <i>s</i>		
ki220%			
	HH ribozyme	binding	site
<:4005 3860			
teqttgtett	tctaggtct		
HI2103 3561			
01115-13			
ON 120 DNA			
-0135 Homo	sapiens		
•**2m2			

HOODS - PONA	HH ribozyme binding site	
(400 + 3861 29tt9tottt	ctaggtctc	19
+310 + 3862 +311 + 19 +312 + DNA +313 + Homo	sapiens	
K.2220 k		
CMM3 - PONA	HH ribozyme binding site	
H400% 3862 attatatatta	taggtctca	19
<pre>&lt;210&gt; 3863 &lt;211&gt; 19 &lt;212&gt; DNA</pre>		
<213> Homo	sapiens	
<220> <223> FCNA	HH ribozyme binding site	
<400 > 3863		
tgtotttota	ggtctcagc	19
<210 · 3864 <211 · 19		
K212 × DNA K213 × Homo	sapiens	
k0000 k k0003 k PCNA	HH ribozyme binding site	
<pre>&lt;400.8 3864 tttotagqtd</pre>	tcagccggt.	19
-00100- 3865 -00110- 19 -00100- DNA		
-1213: Homo	sapiens	
+12205 +12231+ PCNA	HH ribozyme kinding site	
-:400:- 3-65 ictaggtete	ageoggteg	19
	ageoggeeg	± 2
-00100 - 3866 -00110 - 14		

-1212 - DNA -1213 - Homb	sapiens
+120 + +1223 × PONA	HH ribozyme binding site
-(4)0 - 3∀66 	at og aga og
toagooggto	gccgsgacg
+1710 + 3867 +1211 + 1⊎	
+:212 + DNA +:215 + Homo	sapiens
H220 ×	
	HH ribozyme binding site
<400 > 3867	
googgtogto	gogalgito
+210 × 3868 <211 × 19	
<pre>&lt;:310 &gt; DNA &lt;:213 &gt; Homo</pre>	saniens
<2.70%	
	HH ribozyme binding site
<400 + 3868	
togogacutt	egeologietic
<210> 3869 <211> 19	
S2125 DNA S2135 Homo	gariana
	Saptens
+0000+ +00030+ PONA	HH ribozyme binding site
- -1450% 3869	
ogogacqtto	gcccgctct
+2100+3870 +22110+19	
-00120- DNA	
SC2175 Homo	sapiens
+02200+ +02230+ PCNA	HH ribozyme binding site
-:400> 3870	

tagacagata	tgaggetee	19
<pre>&lt;210 + 3871 &lt;211 + 19 &lt;212 + DNA</pre>		
<pre><!--213 + Homo</pre--></pre>	sapiens	
4220 + 4223 + PONA	HH ribozyme binding site	
4400 - 3971 totaaggete	ctgaagoog	19
<pre>8210 + 3672 +211 + 19 &lt;212 + DNA</pre>		
<213 Homo	sapiens	
K2205		
<223> PCNA	HH ribozyme binding site	
-:400 > 3872	g at a ga at t	19
geegaaaeta	gotagaott	1.57
4210× 3873		
4211> 19		
-010 - DNA		
+213 × Homo	sapiens	
-0000		
	HH ribezyme binding site	
-:400:- 3873		
aaactagita	gacttteet	19
-:L10:- 3874		
-211-13		
-07178 DNA		
+12130 Homo	sapiens	
-1710:		
CONSTRUCTION FORM	HH ribczyme binding site	
<pre>&lt;(400) 3874</pre>		
agot agaidt t	tectecte	19
-1210:- 3HT5		
101111 14		
1.1.1 DNA		
H2152 Home	sapiens	

+1000 + +1023 + PONA	HH ribozyme binding site	
<pre></pre> <pre></pre> <pre></pre> <pre>gotagacttt</pre>		19
H210 + 3876 H211 + 19 H212 + DNA		
-1213 - Homo	sapiens	
-(220)- -(225)- PCNA	HH ribozyme binding site	
<pre>&lt;(400 + 3876 ctagactttc</pre>		19
<pre>&lt;210&gt; 3877 &lt;211&gt; 19 &lt;212&gt; DNA</pre>		
<213 > Homo	sapiens	
<2005 <2005 PCNA	HH riboz; me binding site	
<400> 3877 quotitosto	attasagas	19
<pre>&lt;010&gt; 3878 &lt;011&gt; 19 &lt;0105 DUA</pre>		
<2135 Homo	sapiens	
*(2200) *(223) FONA	HH ribozyme binding site	
H4000 3578 titleatesti	ceegeetge	19
-0010:-3879 -0011:-19 -0010:-DMA		
+22130 Homo	sapiens	
	HH ribozyme binding site	
ku40mik 3k79 tilaaraatta	cagaatgaa	19
4210× 3380		

H211 + 19 H212 + DNA H213 + Homo	sapiens	
H2200 H223 + PONA	HH ribozyme binding site	
0400 + 3880 obtgootgta		19
<pre>&lt;210 &gt; 3881 &lt;011 &gt; 19 &lt;010 + DMA</pre>		
<013 × Homo <020 ×	sapiens	
<pre>&lt;0.03 &gt; PCNA &lt;400 &gt; 3881</pre>	HH ribozyme binding site	
tagoggogtt <210 - 3882	gocasteeg	19
<211 - 19 <212 - DNA <213 - Home		
K220 ·	HH ribozyme binding site	
<400 + 3882		19
gttgccactc -0210> 3883 -0211> 19	- byccacca:	1.3
+:213: Homo	sapiens	
-02200- -02235 FCNA	HH ribozyme binding site	
::400:- 3883 ccaccatgtt	cgaggcgcg	19
<1105 3884 <1115 19		
HORITH DNA HORITH Homo	sapiens	
+:220:- +:223:- PCNA	HH ribozyme binding site	

-:400 - 3934 pappatgito	gaggegege	19
#210 + 3#85 #211 + 19 #212 + DNA #213 + Homo	sapiens	
+0000 + +0223 + PONA	HH ribozyme binding site	
::400 - 3885 gogontgato	cagggetee	19
<pre>&lt;010 + 3886 &lt;011&gt; 19 &lt;012&gt; DNA &lt;013&gt; Homo</pre>	sapiens	
K220>	HH ribozyme binding site	
<400> 3886 tecagggete	catoctcaa	19
<pre> 210 &gt; 3587  211 &gt; 19  212 &gt; DNA  213 &gt; Homo </pre>	sapiens	
K000 + K003 + F0NA	HH ribozyme binding site	
k400 + 3887 gggetecate	ctcaagaag	19
0210 - 3888 0211 > 19 0212 - DNA 0213 - Homo	sapiens	
-02001-	HH ribozyme kinding site	
-14001- 3888 chodatests	aagaaggtg	19
:0100 3489 :0115 19 :0.175 DNA :02135 Homo	sapiens	

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42200 42220 PCNA	HH ribozyme binding site	
<ul><li>400 + 3389</li><li>agaaaggtqtt</li></ul>	ggaggcact	19
<pre>&lt;210 + 3890 &lt;211 + 19 &lt;212 + DNA &lt;113 + Home</pre>	sanjars	
<12.10 ·	Sapiens	
	HH ribozyme binding site	
<400> 3890 ggaggcaete	aaggacete	19
<210> 3891 <211> 19 <212> DNA		
<pre>&lt;213&gt; Homo</pre>	sapiens	
+1220 > +1223 > PONA	HH ribezyme binding site	
-400 × 3891 maaggacete	atcaacgag	19
<pre>&lt;210 + 3892 +211 + 19 &lt;212 + DNA</pre>		
<013 - Homo	sapiens	
<220 + <223 + PCNA	HH ribozyme binding site	
<4000 3892 gdaceteate	aacgaggcc	19
00105-3893 00105-19 00105-DNA		
·2135 Homo	sapiens	
+12200+ +12230+ PCNA	HH ribozyme binding site	
-:400:- 3393 tgctgggata	ttageteca	19

+:210 + 3994 +:211 + 13 +:212 + DNA		
-1113 - Homo	sapiens	
-(120 +	HII riboguma binding aita	
	HH ribozyme binding site	
H400 - 3994 Otgaqatatt	agetecage	19
-:210 - 3895		
1211 - 19		
-0112 + DNA -0213 > Homo	sapiens	
K220>		
	HH ribozyme binding site	
<400> 3895		
tqqqatatta	gctccagcg	19
<210> 3896		
<211> 19 <212> DNA		
<213> Homo	sapiens	
SIT 0 >	THE STATE OF THE S	
S35 PUNA	HH ribozyme binding site	
<pre>~400 ~ 3896 atattagete</pre>	cagoggt gt	19
<pre>&lt;210 + 3897</pre> <pre>&lt;211 + 19</pre>		
+0110 + DNA +0110 + Homo	saniens	
-:220		
	HH ribozyme binding site	
+(400) - 3597		
cagoggtata	aacetgrag	19
+01100 - 3A98		
+00110+ 13 +00120+ DNA		
<12131/ Home	sapiens sapiens	
<122(0) ·		
MZZZZZ PCNA:	HH ribozyme binding site	

<pre>&lt;:400 + 3898 graatggactic</pre>	gtcccacgt	19
+:210 + 3899 +:211 + 19 +::12 + DNA		
Hall3% Homo	sapiens	
<220 > < 223 > PONA	HH ribozyme binding site	
+400×3899		
togastogto	ccacgtctc	19
<pre>&lt;310 &gt; 3900 &lt;311 &gt; 19</pre>		
<210 DNA		
<213> Homo	sapiens	
< 0.00 ×		
<223 > PCNA	HH ribozyme binding site	
· 400 · 3900		1.0
gtoccacqto	tctttggtg	19
+ 010 + 3901 + 011 + 19		
· 110 · DNA		
5213 - Homo	sapiens	
· 210 »		
+223% PCNA	HH ribozyme binding site	
400 - 3901		
necaegtete	tttggtjca	19
+2100+3902		
+ 2110+ 19 + 2120+ DNA		
-2130 Homo	sapiens	
- 11 00-		
+1130+ ECNA	HH ribezyme binding site	
· 4000 3 400		
cacqictori	tggtgcagc	19
·0.10> 3903		
C.110: 19		

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(213) Homo	sapiens	
-11.20 + -11.23 + PONA	HH ribozyme binding site	
4400 - 3903 apqueteint		19
-1, 10 + 3904 -1211 - 19		
+:212 + DNA -:213 + Homo	sapiens	
+1220 + +1223> PONA	HH ribozyme binding site	
<pre>&lt;400 x 3904 agtqcagetc</pre>	accetgegg	19
<pre>&lt;210 × 3905 &lt;211 &gt; 19 &lt;212 &gt; DNA</pre>		
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	HH ribozyme binding site	
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H210 > 3906 H211 > 19 H212 > DNA		
<pre>&lt;1213 &gt; Homo </pre>	sapiens	
	HH ribozyme binding site	
ctgagggatt	cgacaccta	19
H2105 3907 K2115 19 KM125 DNA		
<pre>\$401.30 Homo \$401.00</pre>	sapiens	
+200× 3307	HH ribozyme binding site	
tgagggcttc	gacacctac	19

+0010 + 3908 +0011 + 13 +0012 + DNA		
-1213 + Homo	sapiens	
+1120 + +1223 + PCNA	HH ribozyme binding site	
(400)/ 3903 togadadeta	ccgctgcga	19
+310 × 3909 +311 + 19 +313 + DNA		
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	HH ribozyme binding site	
<400 - 3909 ogtgaacots	accagtatg	19
<210> 3910 <211> 19 <212> DNA		
<213 > Homo	sapiens .	
+:220 > +:223 > FCNA	HH ribozyme binding site	
+1400.+ 3910 otoaccagta	tgtccaaaa	19
<pre>&lt;0.105 3911 &lt;0.110 19 &lt;0.110 DNA</pre>		
<pre>*:2170 Homo</pre>	sapiens	
HODDON HUDBEN FONA	HH ribczyme binding site	
<pre></pre>	caaaatact	19
+00100+ 3412 +00100+ 14 +00100+ DNA +00130+ Homo	sapiens	
-:2205		

<pre><!--223 + PINA</pre--></pre>	HH ribozyme bi	nding site		
<pre><i400 +="" 3912="" gtodaaaata<="" pre=""></i400></pre>	ctaaaatgc		1	9
K210 - 3913 K211 - 19 K212 - DNA				
<pre></pre> <pre>####################################</pre>	sapiens			
<pre><d200+ <d223+="" fcna<="" pre=""></d200+></pre>	HH ribozyme bir	nding site		
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<4005 3914 aatgaagata	tcattacac		1	9
H210% 3915 H211% 19 H212% DNA				
<213 > Homo	sapiens			
KIDON KIDBN PONA	HH ribozyme bir	nding site		
::400 × 3915 tgaagatato	attacacta		1	g
+010+ 3916 +0115 19 +010+ DNA				
-17133- Homo	sapiens			
+12200+ +12230+ FCNA	HH ribozyme bir	nding site		
44000 3416				
aga*atcatt	acactaagg		1	9
K210 - 3317				
<211> 19				

-1212 > LENA		
+213 → H⊖mo	sapiens	
·:220 *		
<223 ← PCNA	HH ribozyme binding site	
-:400 × 3917		
gatatoatta	cactaaggg	19
+1210 + 3918 +1211 + 19		
HILLS INA		
-1213 - Homo	sapiens	
: 5: 5:5		
<pre></pre> <pre><pre><pre><pre><pre><pre><pre>&lt;</pre></pre></pre></pre></pre></pre></pre>	HH ribozyme binding site	
2237 13411	mi libonyme binding bice	
<400> 3918		
dattadadta	agggccgaa	19
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KI220 A		
+1223> PONA	HH ribozyme binding site	
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googaagata	acgcggata	19
<210 × 3920		
<2115 19		
<pre>&lt;20108 DNA &lt;2013 - Homo</pre>	saniens	
Trenter	ouplend	
-12200-		
RIZIBN FONA	HH ribozyme binding site	
-:4005 3920		
aacqcggata	cattggaga	19
-m105 3921 -m115 19		
HILLS DNA		
-11135 Homo	sapiens	
-: <u>20</u> 0e-		
	HH ribozyme binding site	
-:4 (no.) - 3 -:01		

oggatacett	ggcgctagt	19
<pre><ul><li>0.10 + 3932</li><li>0.211 + 14</li><li>0.212 + DNA</li></ul></pre>		
+1113→ Homo	sapiens	
00200 0023 + FONA	HH ribozyme binding site	
<pre>&lt;400 × 3922 citqqcqcta</pre>	gtatttgaa	19
0010 + 3923 0011 × 19 0012 > FMA		
+1213> Homo	sapiens	
<220 > 1238 PCNA	HH ribozyme binding site	
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<pre>&lt;010 &gt; 3924 &lt;011 &gt; 19 &lt;0212 &gt; DNA</pre>		
<2135 Homo	sapiens	
H220 × H223 × PCNA	HH ribozyme binding site	
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+02100 3925 +02110 19 +02120 ENA		
+12131+ Home	sapiens	
+01000 +01130+ FCNA	HH ribozyme binding site	
:4000 3925 qotaqtatti	gaagcacca	19
::21(:- 3426 ::211:- 14 ::212:- DNA		
+12135 Home	canions	

-:220 ·		
-11.13 - PONA	HH ribozyme binding site	
-: <b>4</b> 00 + 3926		
qqagaaagtt	tcagactat	19
+1110 × 3927		
0.11 + 19		
-1212 DNA		
+213 > H⊕mo	sapiens	
R120 ×		
	HH ribozyme binding site	
.225 2 3111	In Thompson binding bice	
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qaqaaagttt	cagactatg	19
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<211> 19		
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agaaagtttc	agactatga	19
agaaag coos	agaccacga	
k210 × 3929		
<211 + 19		
4212 - DNA		
<213 - Homo	sapiens	
SILIQ +		
SILIB - PONA	HH ribozyme binding site	
3929 ********	h a a a a h a a a	10
tttdagadta	tgaaatyaa	19
km10 + 3930		
<111 • 19		
-C110 - DNA		
4113 Homo	sapiens	
+17 71c. •		
123 PCNA	HH ribozyme binding site	
H400-3930		
adatqaaqtt	gatggattt	19
√21ar₂ 3.431		
- 2 111 1 5 1 5 L		

H211: 13 H212: DNA H313: H5m5	sapiens	
+0000 + +000A	HH ribozyme binding site	
√400 + 3931 ttgatggatt	tagatgttg	19
+0210 + 3932 +0011 + 19 +0012 + DDA +0013 + Homo	sapiens	
<2200 × +2203 * FCNA	HH ribozyme binding site	
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+:2003 +:003:- FCNA	HH ribozyme binding site	
::400:- 3934 *ttagatgtt	gaacaactt	19
+0108 3935 +0118 19 +0108 DNA +0138 Homo	sanione	
::220:	HH ribozyme binding site	

-:400 - 3935 tgaadaastt	ggaattoca	19
<pre>&lt;210 + 3936 &lt;211 + 19 &lt;212 + DNA</pre>		
<pre>&lt;213 + Homo</pre>	sapiens	
	HH ribozyme binding site	
<pre>&lt;400 - 3936 acttqgaatt</pre>		19
<pre>&lt;210&gt; 3937 &lt;211&gt; 19 &lt;212&gt; DNA &lt;212&gt; DNA </pre>		
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	HH ribozyme binding site	
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<010> 3938 +011> 19 <0108 DMA		
<pre><d13> Homo</d13></pre>	sar·iens	
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44005-3938 aacaggagta	cagctgtgt	19
-1110> 3939 -111> 19 -1110> ENA		
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+2205 H2235 PCNA	HH ribozyme binding site	
<pre>-:40ms 3 439 cagetgtata</pre>	gtaaagatg	19
HI 105 3 440 HI 115 1 4 HI 125 DNA		
-1213> Homo	sapiens	

+:220 + +:2213 + PCNA	HH ribozyme binding site	
<400 + 3940 stgtgtagta	aagatgcct	19
+:210 + 3941 +:211 + 19 +:212 + DNA		
-:213 = Homo	sapiens	
-:010 -	III wilesawa himdina aita	
	HH ribozyme binding site	
K400× 3941 aagatgoott	ctggtgaat	19
<211> 3942 <211> 19 <212> DMA		
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KU205 K2235 PCNA	HH ribozyme binding site	
k4000 3943 otgqtgaabt	tgcacgtat	19
-11101- 3944 -12111- 19		
KUILU DNA KUILU HOMO	sapiens	
+10200+ 	HH ribozyme binding site	
	nn fibozyme binding Site	
<pre><!--400:- 3944 tggtgaattt</pre--></pre>	gcacgtata	19

HC10 + 3945 HC11 + 19 HC12 + DNA HC13 + Homo	saniens	
K1220 +	HH ribozyme binding site	
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8210 + 3946 8211 + 19 8212 + DNA 8213 + Home	sapiens	
<220 <i>0</i> ×	HH ribozyme binding site	
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.(2208	HH ribozyme binding site	
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+0105 3948 +0115 14 +0115 DNA +0135 Homo	sapiens	
+:2205 +:2235 FCNA	HH ribozyme binding site	
H4001-3948 degagatete	agccatatt	19
+0110+ 3949 +0110+ 19 +0011+ DMA +0013+ Romo	sapiens	
+:220:+ +:223:+ PCNA	HH ribozyme binding site	

	ttggagatg	19
2232400110		
+:210 + 3950		
H211 × 19 H212 > DNA		
-1213 Homo	sapiens	
+:220>		
HODB3≠ FONA	HH ribozyme binding site	
-:400> 3950		
daqodatatt	ggagatgct	19
.515 557		
+:210 × 3951 +:211 > 19		
<212> DNA		
-1113> Homo	sapiens	
+:220>		
	HH ribozyme binding site	
<400 × 3951	gtaattaa	19
agatgctgtt	gtaattico	19
<1210 × 3952		
H211 > 19 H212 > DNA		
-1212 - EnA -1213 - Homo	sapiens	
	•	
4(220) >	III wikanan kindina sika	
+DII3 > FUNA	HH ribozyme binding site	
<400 > 3952		
tgotgttata	atttectgt	19
√210 ÷ 3953		
1.12 19		
HITTO ENA		
CDIBO Homo	sapiens	
· Dis		
-CORR PONA	HH ribozyme binding site	
44005 3953		
tatiataatt	tectgtgea	19
<pre>&lt;0.10:- 3954</pre>		
+12111+ 1# +12121+ DNA		
-		

-:213 - Homb	sapiens	
- <u></u>		
	HH ribozyme binding site	
<400× 3954		
gitqtaattt	cctgtgcaa	19
4210 - 3955		
HIII - 19		
<pre><pre></pre><pre><pre>DDA</pre></pre></pre>		
+213 → Homb	sapiens	
H2200		
	HH ribozyme binding site	
	11031 <sub>1</sub> 2110111 <sub>3</sub> 0100	
<400 → 3955		
ttgtaatito	ctgtgcaaa	19
<210> 3956 <211: 19		
KITTA TA		
+213 + Homo	sapiens	
	•	
K220>		
K223≯ FCNA	HH ribozyme binding site	
-:400.8 3956		
gagtgaaatt	ttctqcaaq	19
,-,-,-		
1100-3957		
·111> 19		
<21105 DNA -		
<2130 Homo	sapiens	
k2200+		
HIZIBH FONA	HH ribozyme binding site	
44000 3957		
aqtdaaattt	totgcaagt	19
H2105 3958		
·211> 19		
HIDITAL ENA		
:2135 Homo	sapiens	
-: <u>1112</u> 713		
	HH ribozyme binding site	
.22 - F 1.M	nn libobyme binding site	
-14001-3358		
gtgaaatttt	ctgcaagtg	19

-:210 + 3959 -:211 + 14		
0012 + DNA 0013 + Homb	sapiens	
-1.20 -		
HIDRS - PONA	HH ribozyme binding site	
R400 + 3954 fgaaattitto		1
-0210 + 3960 -0211 + 19		
ILIL DNA		
<213≠ Homo	sapiens	
<220>	HH ribozyme binding site	
√400 - 3960 tgqaqaactt		]
	994440994	
<210 × 3961 <211 × 19		
K212× DNA		
+:213 + Home	sapiens	
+0000 +	III nibawa binding sita	
Control of Editory	HH ribozyme binding site	
<pre>&lt;400 &gt; 3961 tggaaacatt</pre>	aaattotoa	1
	aaaccgcca	
+:210 + 3962 +:211 + 19		
-12 111 - ENA		
-0113 - Homo	sapiens	
-(220)-	WW wibaguma binding site	
	HH ribozyme binding site	
-04005 3962 ggaaacatta	aattotoac	1
	aacegoode	1
+021(0) 3463 +02110 14		
HILL DNA		
-12150 Homo	sapiens	
·:2201-		

H223 → FCNA	HH ribozyme binding site	
-(400 + 3963 abattaaatt	gtcacagac	19
<pre>&lt;210 + 3964 &lt;211 + 19 &lt;212 + ENA &lt;213 + Home</pre>	sapiens	
-(220 -	HH ribozyme binding site	
<400 · 3964 ttaaattgtc	acagacaag	19
<210 × 3965 <211 × 19 <212 + DNA <213 × Homo	sapiens	
<1200 <3235 FONA	HH ribozyme binding site	
<pre>&lt;400× 3965 cagacaagta</pre>	atgtcgata	19
+210% 3966 +211% 19 +212% DNA +213% Homo	sapiens	
-12008 -12030 FONA	HH ribozyme binding site	
:400:-3966 aagtaatgto	gataaagag	19
+010+ 3967 +011+ 14 +012+ ENA +013+ Homo	sapiens	
0220% 02230 FINA	HH ribozyme binding site	
(400)- 3 m <sup>2</sup> aatgrogata	aagaqgagg	19
<1100 3968 <0110 19		

1212 - ENA		
4213 + Homo	sapiens	
-1230 x	THE ACTION AND THE CONTRACT OF	
HUMBA PUNA	HH ribozyme binding site	
-:400 - 3968		
ggaagctgtt	accatagag	19
-210 - 3969		
<211 + 19 ·		
HALLO - DNA		
+1213 + Homo	sapiens	
+:210 >		
	HH ribozyme binding site	
.EED7 TIME	in Tipozyme binding siec	
400 - 3969		
gaagotgtta	ccatagaga	19
H210/ 3970		
<211 • 19		
KIRTON DNA		
<213 - Hemo	sapiens	
-0.20 s		
	HH ribozyme binding site	
<400 + 3970		
tgttaccata	gagatgaat	19
1010 0071		
<2105 3971 <2115 19		
-0113 14 -0113 DNA		
-1213 - Homo	saniens	
11.11.0	buplene	
41 <u>22</u> 5 -		
COOK PANA	HH ribozyme binding site	
44(ng) - 3971 -		19
tqaaddaytt	Caactaact	13
-0:105 3972		
0.115 19		
OF 12% DNA		
-0.130 Hemo	sapiens	
·[[]] (1) ·		
HZLUG - PONA	HH ribozyme binding site	
·:400 - 3972		
-01.01.4		

gaaccagttc	aactaastt	19
+010 + 3973 +011 + 19 +0212 + DNA		
-0213 - Homo	sapiens	
+:220 + +:223 + PGNA	HH ribozyme binding site	
<pre>&lt;400 + 3973 agttcaacta</pre>	acttttgca	19
<pre>&lt;210 + 3974 &lt;211 + 19 &lt;212 &gt; DNA</pre>		
<213 · Homo	sapiens	
<220% <223% PCNA	HH ribozyme binding site	
<400> 3974 daadtaadtt	ttgcactga	19
<pre>&lt;010&gt; 3975 &lt;011&gt; 19 &lt;010&gt; DNA</pre>		
·III > DNA ·III > Homo	sapiens	
+0.20 × +0.23 × PCNA	HH ribozyme binding site	
√400 > 3975 aactaacttt	tgcactgag	19
-0010 - 3976 -0011 - 19		
+212 + DNA +213 + Homo	sapiens	
+:220 + +:223:- PCNA	HH ribozyme binding site	
H4000- 3976 actaactitt	gcactgagg	19
+0.100+ 3477 +0.110+ 19 +0.120+ DNA		
-:213: Hemo	saniens	

-:221		
H223+ PCNA	HH ribozyme binding site	
-:400 - 3977		1.0
dadtqagqta	cctgaactt	19
K210 + 3978		
-211 - 19		
HIII - DNA		
+1213 - Homo	sapiens	
H2208		
	HH ribozyme binding site	
.225 15101	in Tibblyme binding Sice	
<400 > 3978		
acctgaactt	ctttacaaa	19
<210 > 3979 <211 > 19		
<pre><pre><pre><pre><pre></pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre< td=""><td></td><td></td></pre<></pre></pre></pre></pre></pre>		
<pre></pre>	sapiens	
	1	
<:220 >		
S223> FCNA	HH ribozyme binding site	
<4005-3979		
- cotgaactto	tttacasaa	19
<u>.</u>		
-:210:- 3980		
+12111 19		
KIZIZIN ENA	apriora	
KD135 Homo	safiens	
<22200 ·		
$\pm (2.2.33)^{2}$ FCNA	HH ribozyme binding site	
+4008 3980		1.0
tgaacttctt	tacaaaagc	19
-:::10:- 3:481		
-11111-14		
·COLOR DNA		
+01130 Hemo	sapiens	
·(220):		
	HH ribozyme binding site	
2 72121		
-04000 3 +81		
qaadttoitt	acaaaagcc	19
H21UP 3982		

.

+:211 + 19 +:212 + DMA +:213 + Homo	sapiens	
01220 + 01223 + PONA	HH ribozyme binding site	
ki400 + 3982 aactitotita		19
<pre>&lt;210 - 3983 &lt;211 - 19 &lt;212 - DNA &lt;213 - Homo</pre>		
H220 × K223≯ PCNA	HH ribozyme binding site	
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<pre>&lt;310 + 3984 &lt;311 + 19 &lt;312 &gt; DNA &lt;313 &gt; Home</pre>	sapier:s	
-1223 - PCNA	HH ribezyme binding site	
k400 = 3984 dadtocaete	tcttcaacg	19
02105 3985 0211 019 0212 0MA 0213 0 Homo	sapier.s	
+.220.5 +:225.+ FCNA	HH ribczyme binding site	
-14000- 3985 otocactotic	ttcaacggt	19
+210+ 3986 +2110+ 19 +2110+ DNA		
+011%+ Bomo +0120+		
-1232 FCNA	HH ribozyme binding site	

		19
-1210 + 3987 -1211 + 19 -1212 > LNA -1213 > Rumo	sapiens	
1220 + 1223 + Fana	HH ribozyme binding site	
0400 + 3987 dadtatotto	aacggtgac	19
H210 + 3988 H211 > 19 H212 > DNA		
<213≥ Homo	sapiens	
<220> <a><a><a><a><a><a><a><a><a><a><a><a><a></a></a></a></a></a></a></a></a></a></a></a></a></a>	HH ribezyme binding site	
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<210 > 3989 <211 + 19 <212 + DMA		
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+1220 + +1223 + FCNA	HH ribozyme binding site	
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acactcaqta	tgtotgoag	19
3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3990 - 3		
<211> DNA		
-02130 Homo	sapiens	
SCHOON HOLDSHIPPINA	HH ribozyme binding site	
R4000 3490		
icadiat șt d	tgcagatgt	19
+0210+ 3 491 +0210+ 19 +0210+ DMA		
:21:0 Homo	sapiens	

H229→		
HUUBH PONA	HH ribozyme binding site	
-:400 × 3991		
tgbagatqta	ccccttgtt	19
H210 × 3992		
-1211 - 19		
-1212 DNA		
-1213 - Himo	sapiens	
+12.10 +		
	HH ribozyme binding site	
220 10.01	in Tibolyme binding Siec	
+400 × 3993		
tgtacccctt	gttgtagag	19
•		
K210 / 3993		
<2110-19		
KDID> ENA		
H213 % Homo	sapiens	
<2220 >		
KUU3 - PCNA	HH ribozyme binding site	
- -:400 - 3393		
aboodttqtt	gtagagtat	19
45555.0041.	geagageae	
₹210 - 3994		
-:211 - 19		
<212.5 DNA		
$\leq\!\!2.13.6~\mathrm{Hom}_{\odot}$	sapiens	
-1220		
HUMBER FUNA	HH ribozyme binding site	
-:400:- 3994		
cottattata	mantataaa	19
own in the contract	gagtataa	10
-0.11.00 - 3.495		
-00110-19		
-00100- DMA		
:11.130- Homo	sapiens	
+17.7.C1+		
HUTER FONA	HH ribozyme binding site	
<4000 3395		
ttgtagagta	taaaattoo	19
ccycayayıa	cadacege	1

-<210 ⋅ 3396					
4011 - 19					
-11.1 · DNA					
Hull3 - Homo	sapiens				
-1 <u>2</u> 20 -					
HUDBS - PONA	HH ribozyme	binding	site		
·(400 - 3396					
gtagagtata	aaattg:gg				19
*1.11 + 1.#					
HUIL + DNA					
<213 → Homo	sapiens				
<2203					
KDDBS FCNA	HH ribozyme	binding	site		
<4005 3997					
qtataaaatt	gcggatatg				19
<210 × 3998					
<0.11 > 19					
<312× DNA					
<pre></pre> <pre>####################################</pre>	sapiens				
+:220x					
	HH ribozyme	binding	site		
K400> 3998					
attgcggata	tgggacact				19
H2105 3999					
SI:115 19					
O 100 ENA					
	annion:				
Sillias Homo	sapiens				
-( <u>2</u> 26)-					
HILLIGH FONA	HH ribozyme	binding	site		
- -(46ac-3499					
	222242242				19
tgggacantt	addatdCld				13
·(210) · 40(0)					
02110-14					
HODILH DNA					
HOTO: Homo	sapiens				
3.1.	F				
+1212 (C)+					
HIZZBR PCNA	HH ribozyme	binding	site		

(400 + 400) Agganactta	aaatactac	19
0210 + 4001 0211 + 19 0212 + DNA 0213 + Homo	sapiens	
<pre><!--220 + <!223 + PONA</pre--></pre>	HH ribozyme binding site	
<400 - 4001 acttaaaata	ctacttqqc	19
<010 % 4000 <011 % 19 <012 % DNA <013 % Homb	saniens	
<220 s	Sapiens	
R223 + PCNA	HH ribozyme binding site	
<400×4001 taaaatacta	cttggctoc	19
<pre>&lt;:210 &gt; 4003 &lt;:211 &gt; 19 &lt;:212 &gt; DNA</pre>	·	
+1213 + Homo	sapiens	
+220% +223% PCNA	HH ribozyme binding site	
<pre></pre>	ggotoccaa	19
+02100+ 4004 +0011+ 13 +00100+ DNA		
+22130 Homo	sapiens	
KODOOH KODOOH POWA	HH ribozyme binding site	
-(400)- 4004 Nactiggoto	ccaagatcg	19
*12.10% 4005 *12.11% 1 + *12.12% DNA		

-0113 → Homo	sapiens	
+1220 > +1223 > FONA	HH ribozyme binding site	
<pre>&lt;400 &gt; 4005 toppaagate</pre>	gaggatgaa	19
<pre>&lt;210 × 4006 &lt;211 &gt; 19 &lt;212 × DNA</pre>		
<213 > Homo	sapiens	
<pre>+DD0 ≠ *DD3 ≠ PCNA</pre>	HH ribozyme binding site	
<400 × 4006 aaqaaggato	ttaggcatt	19
+210 + 4007 +211 + 19 +212 + DNA		
+:213 + Homo +:2205	sapiens	
	HH ribozyme binding site	
-:400 × 4007 gaaggatott	aggcattct	19
H210 + 4008 H211 > 19 H212 > DNA		
-:113 - Homo	sapiens	
+:220.5 +:223:- PCNA	HH ribozyme binding site	
-(4000- 4008 aaqqatetta	ggcattett	19
+00105 4009 +00105 19 +00105 DNA		
-00130 Bomo	sapiens	
KO1 60+ K21 30+ PONA	HH ribozyme binding site	
<400> 4009 cttaggcatt	cttaaaaatt	19

<pre>#310 + 4010 #311 + 13 #312 + DNA #313 * Homo</pre>	sapiens	
+120 + +122 + + PONA	HH ribozyme binding site	
<pre>&lt;400% 4010 tradgeatts</pre>	ttaaaattc	19
<pre><pre><c10> 4011 <c11> 19 <c11> DNA <c13> Homo</c13></c11></c11></c10></pre></pre>	sapiens	
<220% <223% PCNA	HH ribozyme binding site	
<400 > 4011 aggeattett	aaaattcaa	19
<pre>&lt;010 &gt; 4010 &lt;011 + 19 &lt;012 &gt; DNA &lt;013 + Homo</pre>	sapiens	
+220.+ +223.* PCNA	HH ribozyme binding site	
<400> 4012 udcattetta	aaattcaag	19
HO108 4013 HO111 19 HO129 DNA HO139 Homo	sapiens	
-07000+ -07030+ PCNA	HH ribozyme binding site	
-:400:-4013 tottaaaatt	caagaaaat	19
::10: 40:4 ::11: 19 ::11: DNA ::21:: Homo	sapiens	
10000		

-:220>

HEER PONA	HH ribozyme binding site	
<400 ← 4014 ottagaatto	aagaaaata	19
HU 10 + 4015 HU 11 + 19 HUILDH DNA		
<213 ← Homo	sapiens	
+1020 > +1023 > PONA	HH ribozyme binding site	
k4000 4015 caaqaaaata	aaactaagc	19
<pre><di10> 4016 <di11> 19 <di12> INA</di12></di11></di10></pre>		
<215> Homb	sapiers	
<:220> <:223> FCNA	HH ribozyme binding site	
<pre>&lt;400&gt; 4016 aataaaacta</pre>	agctctttg	19
KC10> 4017 KC11+ 19 KC13> DNA		
SIII 308 Homo	sapiens	
40200+ 40233+ FONA	HH ribozyme binding site	
(400)- 4017 aactaagsto	tttgagaac	19
00100-4113 00110-13 00100-DNA		
-17 130 Homo	sapiens	
002009 00230 FCNA	HH ribozyme binding site	
04005 4018 chaagetett	tgagaactg	19
00100 4019 02112 13		

HOIDH DNA		
4213 + Homo	sapiens	
K129 +		
HIDB + PONA	HH ribozyme binding site	
(4)) ) + 4019		
taayototti	gagaactgo	19
1010 AC 00		
<pre>&lt;0210 * 4020</pre>		
5011 + 19 5011 + 50A		
-1113 - Homo	sariens	
16.2.46	5dp 10115	
K220 ×		
	HH ribozyme binding site	
<400> 4010		
agaactgott	ctaagatgc	19
<210 ~ 4021		
<pre>&lt;211&gt; 19</pre>		
<pre>&lt;212 &gt; DNA   </pre>		
<213 ← Homo	sapiens	
+:210+		
	HH ribozyme binding site	
2223 23111	In The Straing Street	
-:400 + 4021		
gaactgette	taagatgcc	1 5
-		
<210 % 4022		
KI11: 1:		
HIJIB - DNA -		
0113 - Homo	sapiens	
ero men		
-(220 - -(220 -	HH ribozyme binding site	
Sanor Puna	nn libbzyme binding site	
0400× 4022		
actgottota.	agatgocag	19
02108 4023		
12111 1)		
·IIII DNA		
4139 Homo	sapiens	
·021.00		
422.30 PCNA	HH ribozyme binding site	
4400≥ 4023		

tqccagcata	tactgaagt	19
+:210 + 4024 +:211 + 19 +:212 + DNA +:213 + Homo	sapiens	
-1220 >		
+003 × PCNA	HH ribozyme binding site	
-400 × 4004 ccagcatata	ctgaagtct	19
+210 + 4025 +211 + 19		
<212> DNA <213> Homo	sapiens	
<220 >		
<223 > PCNA	HH ribozyme binding site	
<400 × 4025 tactgaagto	ttttctgtc	19
+210 + 4026 +211 + 13		
+ 012 + DNA + 013 + Homo	sapiens	
<.2205		
<2003 → PCNA	HH ribozyme binding site	
.400 × 4006		
otgaagtott	ttotgtoac	19
+210 + 4027 +211 + 19 +212 + DNA		
-1130- Homo	sapiens	
+ 2200+ + 2230+ PCNA	HH ribozyme binding site	
- 4000 - 4017		
tqaaqtottt	totgtoaco	19
<pre>&lt;1100 4028 </pre>		
K2170: DNA K2130: Home	caniane	

-:220-		
4223 - PONA	HH ribozyme binding site	
-(4 10) - 4028		
gaagtotttt	ctgtcacca	19
+(210 + 4029		
+311 - 19		
-1212 - DNA		
$<213 + \text{Hom}\varphi$	sapiens	
<220 ·		
HULB - PUNA	HH ribozyme binding site	
-:400× 4029		
aagtetttte	tgtcaccaa	19
-		
<210 > 4030		
<211> 19		
<212> DNA <213> Homo	ganiang	
%215 * Homo	sapiens	
K220%		
	HH ribozyme binding site	
+:400 × 4030		1.0
cttttctqtc	accaaattt	19
-U10 + 4031	•	
-2211 - 19		
312 DNA		
<213> Homo	sapiens	
-(220's		
	HH ribozyme binding site	
SEE TIME	In Tibolyme binding bice	
-(400 - 4031		
tcaccaaatt	tgtacctct	19
-0110 - 4032 -0211 - 19		
-DIA		
· 213 · Homo	sapiens	
· 220.		
· BH3· PCNA	HH ribozyme binding site	
+ 40m+ 4m32		
dacdaaa*.tt	gtaceteta	19
-02100 4033		

<pre>4211 * 19 4212 * DNA 4213 * Homo</pre>	sapiens	
<000 + + + + + + + + + + + + + + + + + +	HH ribozyme binding site	
H400 + 4033 baaatttqta	cctctaagt	19
+010 + 4034 +011 + 19 +010 + DNA		
<pre>&lt;1.13 + Homo &lt;1.20 &gt;</pre>		
<pre>&lt;400 + 4034 tttgtacete</pre>	HH ribozyme binding site	19
<210 + 4035 <211 + 19 <212 > ENA	taagtacat	1.5
<pre><ill3> Homo</ill3></pre>	sapiens	
	HH ribozyme binding site	
-3400 + 4035 tgtacctcta	agtacatat	19
<pre>%210 % 4036 %001 % 14 %0017 % DNA</pre>		
<pre>+:220's</pre>		
H223H FCNA H400H 4036	HH ribozyme binding site	
cototaaqta	catatgtag	19
:00110 14 :00110 DNA :00130 Homo	sapiens	
-12201+	HH ribozyme binding site	

-(40 )> 4037 taagtabata		19
H010 + 4038 H011 + 13 H012 + DNA H013 + H0mp	sapiens	
<pre><pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre><td>HH ribozyme binding site</td><td></td></pre></pre></pre></pre></pre>	HH ribozyme binding site	
-(400 × 4038 tabatatuta	gatattgtt	19
<210 > 4039 <211 > 19 <212 > DNA		
<213> Homo	sapiens	
+:220 > -:223 > FONA	HH ribozyme binding site	
<400> 4039		
tatqtaqata	ttgttttct	19
+110> 4040 +111> 19 +111> DNA		
+1213> Homo	sapiens	
<2200 × <223> FONA	HH ribozyme binding site	
<pre>&lt;400% 4040 tqtaqatatt</pre>	gttttetgt	19
+02108 4041 +02118 19 +02109 DNA		
+12130 Homo	sapiens	
+12200 +		
HULBS PANA	HH ribozyme binding site	
<pre>&lt;4000 4041 adatatt@tt</pre>	ttctqtaaa	19
*C10: 4:42 *C11: 1 *		
<pre><c1200 <="" dna="" pre=""></c1200></pre>	sapiens	

4230 + 4233 ← PONA	HH ribozyme binding site	
<pre>&lt;:400 &gt; 4042 gatattgttt</pre>		19
HILLON 4043 HILLON 14 HILLON DNA		
<pre>************************************</pre>	sapiens	
	HH ribezyme binding site	
<400> 4043 atantgtitt		19
<pre><d210> 4044 <d211> 19 <d212> DMA <d213> Hemo</d213></d212></d211></d210></pre>	sapiens	
+:220 ×	HH ribozyme binding site	
<400 × 4044 tattottto	tgtaaataa	19
<pre>*1210 + 4045 *1211 &gt; 19 *1212 &gt; DNA *1213 &gt; Homo</pre>	sapier.s	
KOZUG KOZEG PONA	HH ribczyme binding site	
K400% 4045 giittotgia	aataaccta	19
+00100+ 4046 +00110+ 19 +00120+ DNA +02130+ Homo	sapiens	
·(2 <u>1</u> (c)·	HH ribozyme binding site	
<400> 4046 totgtaaata	acctatttt	19

<pre>&lt;310 &gt; 4047 &lt;311 &gt; 19 &lt;3212 &gt; DNA &lt;313 &gt; Homo</pre>	anniona	
7.1.137 n.mo	sapiens	
-1220 - -1223 - PONA	HH ribozyme binding site	
<pre>&lt;400&gt; 4047 aaataacota</pre>	tttttttc	19
H210 × 4048 H211 > 19 H211 > DNA H213 > Homo	saniens	
+:220> +:223> PONA	HH ribozyme binding site	
<400> 4048 ataacctatt	ttttttctc	19
<pre>#210&gt; 4049 #211&gt; 19 #212&gt; DNA #213&gt; Homo</pre>	sapiens	
HAAOS HAAAS PONA	HH ribozyme binding site	
<400 × 4049 taacctaitt	tttttctct	19
H2108 4050 H2118 19 H2128 DNA H2138 H0mo	sapiens	
+:200:+ +:203:+ PCNA	HH ribczyme binding site	
-:400:- 4080 aaoctatitt	ttttctcta	19
:0.100: 4051 :0110: 19 :0110: DNA :0130: Homo	sapiens	
<22000 <22000 PCNA	HH ribozvme binding site	

400 → 4051 abotatttt	tttctctat	19
H210 + 4052 H211 + 1 + H212 + DNA H213 + H5mb	sapiens	
+1120 + +1123 + P€NA	HH ribozyme binding site	
<pre></pre>		19
<pre>&lt;210 &gt; 4053 &lt;211 • 19</pre>		
<012 > DNA < 0213 > Homo	sapiens	
+12205 +123> PCNA	HH ribozyme binding site	
<pre>&lt;400 + 4053 ctattttttt</pre>	tctctattc	19
-210 - 4054 -0011 - 13 -0212 - DNA		
-0213 × Homo	sapiens	
	HH ribozyme binding site	
tattttttt:	ctctattct	19
-0.110 19 -0.110 00A -0.1130 Homo	ganiona	
+:2.2.00+		
-(400) 40€5	HH ribozyme binding site	10
arttittita <1.10: 4056	tctattctc	19
<pre><ul><li>(211): 1 )</li><li>(212): DNA</li></ul></pre>		

-:213 → Homo	sapiens		
-1203 + PONA	. HH ribozyme binding site		
-:400 - 4056 httptttchc			19
+010 + 4057 +011 + 19 +011 + DNA			
+1213 - Homo	sapiens		
+:220 + +:223 + PCNA	. HH ribozyme binding site		
<400% 40%7 tttttctcta			19
<210 > 4058 <211 > 19 <211 > DNA			
<213 · Hemo	sapiens		
<220 > HUD3 - PCNA	HH ribozyme kinding site		
H400 + 4058 titototati		•	19
+0010 + 4059 +00115 19 +00115 DNA			
-1213 · Homo	sapiens		
+120 e + +110 3 > PONA	HH ribozyme binding site		
<pre>&lt;400&gt; 4059 thototatto</pre>		:	19
+0210 + 4060 +0211 + 19 +0212 + DNA			
+12131 Hamo	sapiens		
+1220 + +12237 PONA	HH ribozyme binding site		
<400 + 4060 ctctattctc	tccaatttg	;	19

<pre>&lt;2105 4061 &lt;2211 - 19 &lt;2212 - DNA &lt;2213 - Homp</pre>	sapiens	
-0000 + -0003 + PONA	HH ribozyme binding site	
<pre>&lt;400.0 4061 etattotete</pre>	caatttgtt	19
H210 + 4062 H211H 19 H212 + DNA K213 H0mo	sapiens	
<220× <223× PCNA	HH ribezyme binding site	
0400 = 4062 Stotocaatt	tgtttaaag	19
HIIO> 4063 HIII> 19 HIII> DNA		
+0135 Hemo	sapiens	
	HH ribozyme binding site	
<pre>&lt;400 &gt; 4063 totocaattt</pre>	gtttaaaga	19
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	sanjens	
·10.00.	HH ribozyme binding site	
H4061- 4064 ccaatttatt	taaagaata	19
+00100+ 4065 +0010+ 19 +0010+ DNA		
Homo	sapiens	

H223 ← PCNA	HH ribozyme bin	ding site		
-:400 + 4065 -:aastitgstb	aaagaataa		1	9
HU10 + 4066 HU11 + 19 H212 + DNA				
<pre></pre> <pre><pre># A Tombook </pre></pre>	sapier.s			
<pre>&lt;:020 + :0203 + FONA</pre>	HH ribozyme bin	ding site		
<400 - 4066 aatttgitta	aagaataaa		1	. 9
<210 - 4067 <211- 19				
<213> DNA <213> Homo	sapiens			
K220> K223> FCNA	HH ribozyme bine	ding site		
<400> 4067				_
ttaaagaata	aagtocaaa		1	9
<210 % 4068 <211 % 19				
H2125 ENA H2135 Homo	sapiens			
<0205 <0205 FONA	HH ribozyme bind	ding site		
-:4005- 4065 gaataaagto	annatata		1	9
gaa .aaaguu -22125 4069	cadageseq		1	J
:00115 19				
-011.08 EMA -02138 Homo	sapiens			
-02105 -02135 PCNA	HH ribozyme bind	ding site		
·:400:- 4569				
gtocaaagto	tgatctggt		1	g
+12101+ 4170 +1211> 13				

<21.5 DNA <313 Homo	sapiens			
+1020 + +1023 + PONA	HH ribozyme	binding	site	
k(400 + 4070 aaytotgato	tggtctagt			19
<pre>&lt;210 + 4071 &lt;211 + 19 &lt;310 + 5NA</pre>				
<pre>&lt;013 + Homo </pre>	sapiens			
<pre>&lt;223 + PCNA &lt;4400 &gt; 4071</pre>	HH ribozyme	binding	site	
tgatotggto	tagttaacc			19
<pre>&lt;210 + 4072 &lt;211 + 19 &lt;212 + DNA &lt;213 &gt; Homo</pre>	sapiens			
K220 - K223 - PCNA	HH ribozyme	binding	site	
<400 - 4072 atotqqtota	gttaaccta			19
<pre>8210 + 4073 82110 19 82120 DNA 82120 Home</pre>	sapiens			
+00000 +00030 PONA	HH ribczyme	binding	site	
c400 4973 tggtdtagti	aacctagaa			19
+:210:- 4074 +:211:- 19 +:212:- DNA -:213:- Homo	sapiens			
-07200- -02230- PCNA	HH ribozyme	binding	site	
·:400> 4074				

ggtstagtta	acctagaag	19
H210 + 4075 H211 + 18 H212 + DNA		
-0013 - Homb	sapiens	
+1220 + +1223 + PONA	HH ribozyme binding site	
-(400 - 4075 aqttaadota	gaagtattt.	19
+0010 + 4076 +0011 + 19 +0010 + DNA		
<213 > Homo	sapiers	
×(220).⊁		
	HH ribozyme binding site	
<400× 4076		19
cetagaaqta	tttttgtct	13
<210 - 4077 <211 > 19		
<2125 DNA		
+:213> Hemo	sapiens	
+12000+ +12030+ PCNA	HH ribozyme binding site	
-:400> 4077		
tagaagtatt	tttgtctct	19
+02100+ 4078 +02110+ 19 +01120+ DNA		
SIN130 Homo	sapiens	
4( <u>2</u> 2 <b>0</b> (+		
	HH ribozyme binding site	
-:400:- 4078		
aqaaqtaftt	ttgtctctt	19
+:210:+ 4079 +:211:+ 19		
<2120 DNA		
<213> Homo	sapiens	

4220 + 4223 + PONA	HH ribozyme binding site	
0400 + 4079 gaagtatitt	tgtctctta	19
+0.10 + 4080 +0.11 + 19 +0.212 + DNA +0.213 + Homb	saniers	
<0.00 e	HH ribozyme binding site	
-:400 - 4080 aagtatttt		19
<2105 4081 <2115 19		
<212> DNA <213> Homo	sapiens	
<pre>&lt;:DD0&gt; <dd3> FCNA</dd3></pre>	HH ribozyme binding site	
<400 × 4081 tatttttatd	tcttagaaa	19
<pre>&lt;010&gt; 4082 &lt;011&gt; 19 &lt;0128 DNA</pre>		
+12135 Homo +12205	sapier.s	
-01230 PCNA -04000 4082	HH ribozyme binding site	
tttttgtbtc +01100-4083	ttagaaata	19
+00110+19 +00100+ DNA +00130+ Homo	sapiens	
·(220)·	HH ribozyme binding site	
-(400)- 4183 thtatothtt		19
.main whow		

.

+:211 + 19 +:212 + DNA +:213 + Homo	sanions	
	Sapiens	
H229 + FONA	HH ribozyme binding site	
-(40)) - 4084		
itgintotta	gaaatactt	19
<pre>&lt;210 - 4085</pre>		
-211 · 13		
+213 + Homo	sapiens	
<220.		
RIDDBH PONA	HH ribezyme binding site	
<400> 4085		1 (
ottaqaaata	cttgtgatt	19
<pre>&lt;210&gt; 4086 &lt;211&gt; 19</pre>		
HOLL FINA		
<2135 Homo	sapiens	
-0.200 %		
FILLUS + FICNA	HH ribozyme binding site	
<400 - 4086		19
aqaaatastt	gigatitit	15
<210 × 4087 <211 × 19		
-1212 · LNA		
213 Homo	sapiens	
-0120 ×		
HILLSH HUNA	HH ribozyme binding site	
-14000 4087	tttataata	19
actidatqatt	tttataata	19
H210H 4088 H211H 19		
-12171: FNA		
+2130 Homo	sapiens	
·:::2(::)		
K223> FCNA	HH ribozyme binding site	

ttataatac	19
sapiens	
HH ribozyme binding site	
tataataca	19
sapiens	
ataatacaa	19
HH ribozyme binding site	
taatacaaa	19
sapiens	
HH ribozyme binding site	
atacaaaag	19
saniens	
	sapiens  HH ribozyme binding site  tataataca  sapiens  HH ribozyme binding site  ataatacaa  sapiens  HH ribozyme binding site  taatacaaa  sapiens  HH ribozyme binding site  taatacaaa

-c221→		
HILLE + PINA	HH ribozyme binding site	
H400 × 4003		
ttttataata		19
·1210 × 4034		
<:211 · 19		
ANG - LIEB		
HDD3 > Homo	sapiens	
H220 +		
	HH ribozyme binding site	
200	III	
<400 - 4094		
caaaagggto	ttgactsta	19
K210 + 4095		
K211 - 19		
:1110 IB :1110 - DNA		
HD13 - Homo	gariera	
Talab t noneo	Sapiter.s	
-1000-		
KODES FONA	HH ribezyme binding site	
<400 × 4095		
	gagtetana	19
aaagggtott	gactetada	1.7
H210× 4096		
-1211 > 14		
SOID> DNA		
<2135 Homo	sapiens	
2.5		
-1110 ×		
ALLISA BUNA	HH ribezyme binding site	
(400) - 4096		
gtottgabto	taaatgcag	19
-11 100 - 40 97		
·0.110 19		
-1111 DNA		
·1130 Homo	sapiens	
+(%)_(0)+		
	HH ribozyme binding site	
MALE OF THE PROPERTY.	in Tibozyme binding Sise	
·:400> 4J97		
cttgactcta	aatgcagtt	19

H210 + 4098 H211 + 19 H212 + DNA		
-0213 - Homo	sapiens	
+1220 + +1223 + PONA	HH ribozyme binding site	
<pre>&lt;400 - 4098</pre> <pre>aaatqcaqtt</pre>	ttaagaagt	19
<pre>&lt;210 + 4099 &lt;211 + 19 &lt;212 + DMA</pre>		
-1213 - Homo	sapiens	
<220 × <223 > PCNA	HH ribozyme binding site	
₹400× 4099 aatqcagttt	taagaagtg	19
+010+4100 +011+19 +012+DNA		
<213> Homo	sapiens	
+0000 + <0003 + PONA	HH ribozyme binding site	
<pre>&lt;400 + 4100 atgcagtttt</pre>	aagaagtgt	19
<pre>&lt;2108 4101 \$0118 19 +0108 ENA +0138 Eomo</pre>	sapien <i>s</i>	
·(2200·	HH ribozyme binding site	
04000 4101 tqcaqtttta	agaagtgtt	19
+00100+ 4102 +00110+ 13 +00100+ DNA		
+1211fth Bumo	sapiens	
+12200+ +1223> PCNA	HH ribozyme binding site	

-:400 → 4102 aagaagtgtt	tttgaattt	19
+:210 + 4103 +:211 + 19		
HONA Homo	sapiens	
<pre><!--200 + </pre--></pre>	HH ribozyme binding site	
+400 + 4103 agaagtgttt	ttgaattta	19
<210 × 4104 <211 + 19		
<212 - DNA <213 - Homo	sapiens	
<223 + PCNA	HH ribozyme binding site	
<pre>&lt;400 &gt; 4104 gaagtgtttt</pre>	tgaatttaa	19
+0010 + 4105 +0011 + 19 +0010 + DNA		
+1113 + Humo	sapiens	
	HH ribozyme binding site	
-:400 - 4105 aagtqttttt	gaatttaaa	19
+:::10 + 4106 +::211 + 13 +::212 + DNA		
<pre></pre> <pre><pre>+:2130 Homo</pre></pre>	sapiens sapiens	
	HH ribozyme binding site	
<pre>%4000 4106 thtttgaatt</pre>	taaataaag	19
<2105 4107 (2115 1) (2125 1914)		

<213> Homo	sapiens	
-0220 + -0225 + PONA	HH ribozyme binding site	
-(400 - 4107 tittgaatit	aaataaagt	19
+010 + 4108 +011 + 19 +012 + DMA		
-1213 - Homo	sapiens	
+:220 + +:223 + PCNA	HH ribozyme binding site	
<pre>&lt;:400 + 4108 tttgaattta</pre>	aataaagtt	19
<pre>&lt;:210 + 4109 &lt;:211 + 19 &lt;:212 + DNA</pre>		
HOTER Home	sapiens	
<pre></pre> <pre><pre></pre> <pre><pre></pre> <pre></pre> <pre><td>HH ribozyme binding site</td><td></td></pre></pre></pre>	HH ribozyme binding site	
<pre>&lt;400 + 4109 autttaaata</pre>	aagttaott	19
+0.100+ 4110 <0.115 19 <0.115 DNA		
-02135 Homo	sapiens	
H228H PONA	HH ribozyme binding site	
-:400:- 4110 aaataaaytt	acttgaatt	19
+0.100 4111 +0.110 10 +0.110 DNA		
-C 130- Homo	sapiens	
-0.100+ -02750+ PCNA	HH ribozyme binding site	
<400> 4111 aataaagtta	cttgaattt	19

```
·:::10 · 4112
\pm 1.11 \pm 19
-1.11 - DNA
< 213 + Homo sapiens
-1230
HIZZB · PONA HH ribozyme binding site
\pm 409 \pm 4112
aaaqttactt gaatttcaa
                                                                              19
\pm 0.010 \pm 4113
\pm 211 \pm 19
·2122 DNA
<213> Homo sapiens
+1220>
<223> PONA HH ribozyme binding site
+4008 4113
tacttgaatt tcaaacaaa
                                                                              19
<d210 > 4114
HD115 19
<\!\!1112\!\times 100A
Hall3 - Homo sapiens
-1220 -
\pm 223 \pm \text{PCNA} HH ribozyme binding site
4400 > 4114
actiquatit caaacaaaa
                                                                              19
\pm 0010 \pm 4115
02225 19
HALLS DUA
\cdot 2130 Home sapiens
400000
HLLL: FONA HH ribozyme binding site
-14000-4115
                                                                              19
cttgaat*tc aaacaaaaa
+00100 411€
-2211: 5.7
HILL DIA
+:213: Homo sapiens
·:400: 4116
```

```
5.2
  gasssctgag aaaggcassa gagaaacaca sttggtggta tattasstgg ta
 :1.10 → 4117
 √211 > 53
 HIZIZ + DNA
 +:213 → Homo sapiens
 -1400 + 4117
                                                                                                                                                                                                                                                                           5.2
  yagodotyay aaaggoacca gagaaacaca ottggtggta tattaootgg ta
 \pm 0.010 \pm 4113
 \pm 1211 \pm 51
 \pm 212 + \text{DNA}
 +1213 · Homo sapiens
 <400≥ 4118
 qagooctqaq aaaggcacca gogaaacaca cttggtggca tattacctgg ta
<210 > 4119
<2111 > 61
 RITTO - DNA
+313 · Homo sapiens
-1400 - 4119
 gagocotgag aaaggoacoa gogaaacaca oggacuucgg toogtggcat attacotggt 60
·1210 · 4120
111 14
1112 DNA
+ 213 Homo sapiens
< 220 >
 <223  General Hairpin Ribozyme Site</pre>
<!!!
<!-- The state of the
KID1 - modified_base
KID1 - (1)
KID3 - Where n is a, c, g or u
\{1,2,3,3,5\}
-0.315 modified base
· 100 (2)
+Tils Where n is a, c, g or u
\{(x,y)\in X_{k+1}^{(k+1)}(x,y)\}
+0.110 mudified base
Rando Where n is a, c, g or u
-121.01-
%221> modified base
```

```
·12.111 \ (5)
H2003 · Where n is a, c, g or u
-1221-
H221 - modified_base
H222 - (9)
H223 - Where n is a, c, g or u
122201 ·
:221 modified_kase
+3.23 \cdot \text{Where n is a, c, g or u}
-1220 -
-221 - modified base
· 2129 (11)
·223 > Where n is a, c, g or u
7220 ×
<221 > modified base
<222 \ (12)
<223 Where n is a, c, g or u
<:220 ×
<221 * modified base</pre>
<2222 (13)
3003. Where n is a, c, g or u
-:2200
+221 - modified base
-1.1.1.1 - (14)
AMMAN Where n is a, c, g or u
-14000 4120
nnibnguenn nnnn
<0.10> 41.11
-00110-15
ACCION DNA
%1135 Homo sapiens
-12.201-
-0.50 General Hairpin Ribozyme Site
-12200
::211: modified_base
::212: (1)
::213: Where n is a, c, g or u
-12.205-
-1121: modified base
-0.1111- (2)
\pm 0.0230 Where n is a, c, g or u
<2200
```

```
HCC1 + modified_base
HCC2 + (3)
HCC23 + Where n is a, c, g or u
·(220) ·
%221 / modified_base
%222 / (5)
\pm 223 . Where n is a, b, g or u
-1220 -
....1 modified base
HDDD (9)
Hills: Where n is a, b, g or u
-1220u -
+:221 · modified base
H222 (10)
<223> Where n is a, c, g or u
<0220 ×
+221 modified_base
<2220 (11)
<223> Where n is a, c, g or u
<0000>
<221> modified base
<2225 (12)
<223> Where n is a, c, g or u
40200 N
SCOIN modified_base
+CLOIN (13)
+CLOIN Where n is a, c, g or u
+ 2200s
<221> modified_base
<2225 (14)
%2230 Where n is a, c, g or u
- _1_-
::1:1: modified base
<1...1 (15)
\times \mathbb{M}(\mathbb{N}) Where n is a, c, g or u
\pm (4\,0\,0) \leq 4\,1\,2\,1
nord-riguerin nnnnn
+111100 - 4122
40010 16
HILLION DNA
1213 Homo sapiens
4:22(C)+
H223> General Hairpin Ribozyme Site
```

```
-:221-
Hintl + modified_base
Hintle (1)
AMM3 · Where n is a, b, g or u
+00.0 >
+00.10 > modified_base
·1. 1.5 • (2)
-0013 · Where n is a, c, g or u
<C210 *
<C211 * modified_hase
<C111 * (3)
<C113 * Where n is a, c, g or u</pre>
\star (11110) +
diff: modified_base
<2220 - (5)
<1.13> Where n is a, c, g or u
<2220>
<221> modified_base
<222 > (9)
<223> Where n is a, c, g or u
7220×
+2210 \times \text{modified base}
+2222 € (10)
32230 Where n is a, c, q or u
\{(j,j),(j,j)\}
+W21+ modified_base
+W222+ (11)
\pm 0.023 \, \cdot \, Where n is a, c, g or u
S0000
-CLD10 modified_base
-CLD10 (12)
-CLD30 Where n is a, c, g or u
+CD08
+CD01: modified_base
+CD00: (13)
+CD03: Where n is a, c, g ir u
Hill: modified_base
Hill: (14)
Millish Where n is a, c, g or u
· [[]][(-]-
+2.2.11 modified base
\pm 0.22 \, \mathrm{M}_{\odot} Where n is a, c, g or u
```

-1220>

```
+3321 \cdot modified base
\pm 2.23 \cdot \text{Where n is a, c, g or u}
+(4)(0) + 4122
nninbiquenn nnnnnn
-:L10 - 4123
\cdot 1.111 \cdot 17
KILLL - DNA
HL13 · Homo sapiens
<12.20 ×
H223 · General Hairpin Ribozyme Site
Hill modified_base
Hill (1)

#2003 - Where n is a, c, g or u
<2200
+321 + m=dified_base
K222% (2)
%223> Where n is a, c, g or u
<12200
<1.21 \times modified_base
\pm 122.8 (3) \pm 122.8 (3) \pm 122.8 Where n is a, c, g or u
+[\underline{-1}]\underline{-1}[1]S
-:221> modified base
·1222> (*)
<223> Where n is a, c, g or u
<0.112
*Miles modified base
*Miles (*)
-III35 Where n is a, c, g or u
- 22068
+1.115 modified base
HR:228 (10)
-00250 Where n is a, c, g or u
+0.1200+
+m215 modified base
+00120+ (11)
+00140+ Where n is a, c, g or u
·::20:-
\cdot 2210 \cdot modified_base
\cdot \mathbb{Z} \, 22 \, \cdot \, \left( \, 12 \, \right)
\pm 223 Where n is a, c, g or u
```

```
-:::2n -
 ::221 + modified_base
::223 + (13)
 +223 Where n is a, c, g or u
 -(220-
 :C21 - modified_base
:C22 - (14)
 8223 Where n is a, c, g or u
 (1)
  \cdot: \mathbb{N} \mathbb{N} + \mathbb{N} = \mathbb{N}  modified base
 \times 222 \times (15) \times 223 \times \text{Where n is a, c, g or u}
 \{(\underline{1},\underline{1},0)_{i\in I}
 <221 \times \texttt{modified} base
 <2222 > (16)
 <\!223\times Where n is a, c, g or u
<:220>
 <221> modified base
 <2225 (17)
 KARAS Where n is a, c, g or u
<400> 4123
 nnnbnguenn nnnnnnn
-12100 4124
 . 211 18
 HILLE DNA
 +2135 Homo sapiens
+12.200+
 <
+12200
+22230 General Hairpin Ribezyme Site
 -17.701-

*D210 mcdified_base

*D210* (1)

*D130* Where n is a, c, g or u

**D130***
**The mcdified_base

**The mcdifi
 \{(\underline{T},\underline{T},\underline{Q})\}_{t=0}^{T}
 \begin{array}{ll} + 12211 + \bmod ified\_base \\ + 12221 + (2) \end{array}
 +2233 Where n is a, c, g or u
-12201-
 \pm 2211 \pm \texttt{modified\_base}
·:2221 (3:
 4223> Where n is a, c, g or u
```

```
>1220>
....l > modified_base
+CLLL* (5) +CLLL* \mbox{ Where n is a, c, g or u}
+(1,20) \leq
dll1 > modified_base
dlll2 > (9)
< 0.033 \times Where n is a, c, g or u
+(\frac{\pi}{2})\frac{\pi}{2}(1)>
H221: modified_base
H222: (10)
H213> Where n is a, c, g or u
<dd0.
<L21> modified_base
<l222 (11)</pre>
<223 Where n is a, c, g or u
<2200
<221> modified base
<222 > (12)
MOD35 Where n is a, c, g or u
<2205
\sim221 \rightarrow modified_base
42220 (13)
\pm 1233 Where n is a, c, g or u
\{1,1,2,0\}
SUBLY modified_base
SUBLY (14)
\leq 2.23 \cdot \text{Where n is a, c, g or u}
<2220>
+221 modified_base
<!2221+ (15)</pre>
\cdot (223)^{2} Where n is a, c, g or u
+00010+ modified_base
+00000+ (16)
\pm 0.1\,\mathrm{M}_\odot Where n is a, c, g or u
+1<u>2.2</u>11.5+
+:2015 modified_base
+:2015 (17)
R223: Where n is a, c, g or u
\cdot \text{CLII} \cdot \text{modified base}
+0.0221 \cdot (18)
32230 Where n is a, c, g or u
```

<400> 4124

nnbnguenn nnnnnnn	18
(210 + 4125)	
+0.211 + 16	
CO12 + DNA	
-0013 - Homo sapiens	
-:220 ·	
<pre>42.23 - Hairpin ribozyme recognition site for cdc 2 kinase</pre>	
<400 + 4125	
	16
acttogtoat coaaat	10
< 4126	
<211 > 16	
K212× DNA	
KIIIAH Homo sapiens	
K220>	
<223% Hairpin ribozyme recognition site for cdc 2 kinase	
<400× 4126	
atatagtoag tottoa	16
acadagicay coccea	10
<210% 4127	
KL117-16	
HILLS + DNA	
<pre>S213 Home sapiens</pre>	
CLO3- Haurpin ribozyme recognition site for cdc 2 kinase	
-:400b-4127	
agtbagtbtt caggat	16
0.108-4128	
-0.110-16	
HOTEL DIA	
<pre><!--!!! Homo sapiens</pre--></pre>	
+122 ob-	
+:2238 Halrpin ribczyme recognition site for odc 2 kinase	
12. Maripin Tibezyme recegnition site for ede 2 kindse	
-04(00)5 410.8	
tootggtmag tacatg	16
-C:10:- 4129	
G111:- 1r	
- COLOR DNA	
<213> Homo sapiens	

Hairpin	ribozyme	recognition	site	for	cdc 2	kinase	
(400 · 4129							
quirtgtcac to	taga						16
-1.10 + 4130							
-1211 - 16							
HUIU - DNA HOMO sap	oiens						
<pre>%020* %003 - Hairpin</pre>	ribozvme	recognition	site	for	ode 2	kinase	
-		g					
<400 · 4130	7a++						16
otggggthag oto	Jytt .						10
<2105							
<2110 <2125 DNA							
<213 Homo sar	piens						
<020 >							
<223>							
<400>							
HARLON 4131 HARLON 16							
SIRILS ENA							
<213> Homo sap	piens						
H22UN							
00033> Hairpin	ribozyme	recognition	site	for	cycli	n Bl	
-:40()> 4131							
tocqagtoac caq	ggaa						16
+:210:- 4132							
1711: 16							
HILLS DNA							
-02135 Homo sap	rens						
+02000+	. ,			_		D.1	
<pre></pre>	ribozyme	recognition	site	tor	cyclin	u BI	
::400: 4132							
dcad*.gtctg ago	ccag						16
32107 4133							

.

```
4211 - 16
HULL DNA
\pm 213 + \text{Homo sapiens}
Hairpin ribozyme recognition site for cyclin B1
\pm 1400 \pm 4135
                                                                    16
ustaigt mag gottto
\pm (210 + 4134)
4211 \pm 16
HILLS - DNA
+2213 · Homo sapiens
-1220 -
+323 · Hairpin ribozyme recognition site for cyclin B1
\pm (400) \approx 4134
                                                                    16
aagbagtbag abbaaa
出2105 4135
\pm 2115 - 16
<2212 > DNA
H213> Homo sapiens
<:2205</pre>
3.235 Hairpin ribozyme recognition site for cyclin B1
+14005 4135
actoggtogg gaagto
                                                                    16
<210> 4136
02115-16
HL112 DNA
+12135 Homo sapiens
<22208
Hairpin ribozyme recognition site for cyclin B1
-400> 413€
tractificate cattat
                                                                    16
<11100 4137
<11110 1+c
+.210> DNA
·21: Homo sapiens
-12201-
<225> Hairpin ribozyme recognition site for cyclin Bl
```

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H:400 × 4137 H:tqqtgtcac tgccat	16
+0210 + 4138 +0211 + 16 +0213 + ONA	
+2213 × Homo sapiens	
<pre>&lt;(220 &gt; </pre> <223 > Hairpin ribozyme recognition site for cyclin B1	
<pre>-:400x 4138</pre>	1 લે
<pre> <pre>       <br <="" td=""/><td></td></pre></pre>	
<pre>&lt;011 &gt; 16 &lt;010 &gt; DNA</pre>	
<pre>&lt;013&gt; Homo sapiens</pre>	
<pre>&lt;:220&gt; &lt;223&gt; Hairpin ribozyme recognition site for cyclin B1</pre>	
<400> 4139	
totgggtogg octoba	16
<210> 4140	
<211 × 16	
HILLS DNA	
+:213> Homo sapiens	
+(220.8)	
<223> Hairpin ribozyme recognition site for cyclin B1	
+:400:+ 4140	
tacctgf rat atactg	16
+1110) + 4141	
+211> 16	
HOLIO FINA	
+:213: Hamo sapiens	
(12.0):	
(223) Hairpin ribozyme recognition site for cyclin Bl	
-:400:4141	
atgtagtidat ggtaaa	16
HC10H 4142	
H2110 16	
82132 ENA 82132 Homo sapiens	

<220→						
			_	2.1	D.1	
-:::::: Hairpin ribozyme	recognition	site	ior	cyclin	RI	
-:400 + 4142						
tyadigidaa gaadaa						16
+(210 + 4143						
<211 × 16						
HARIA DNA						
-0013 - Homo sapiens						
Saprens						
+(220 ×						
Hairpin ribozyme	rocomition	aita	£ ~ ~	DCHA		
nalibin ilpozyme	recognicion	SILE	TOL	PUNA		
<400 > 4143						
						16
gagtygtogt tgtott						16
<210 - 4144						
K2115 16						
KULU + DNA						
+1213 - Home sapiens						
mis nome suprens						
+:220 +						
+323 · Hairpin ribozyme	recognition	eita	for	תוזים		
nalipin libozyme	recognition	2166	T.OT	LONA		
-:400 - 4144						
togutgtott totagg						16
Jugungener codage						11)
+1110 + 4145						
HŽ11.+ 16						
*2125 DNA						
+2130 Home sapiens						
•						
*::20°						
+22233 Hairpin ribozyme	recognition	site	for	PCNA		
	-					
<14.000 - 4145						
geetagtica gggete						16
100 1 20 S						
H210% 4146						
K(211) - 16						
+:212:- DNA						
+C213> Homo sapiens						
+:2200						
			c	Dans		
Hairpin ribozyme	recognition	site	ior	PUNA		
.:10m: 4146						

gastogtods adgitet	16
+1010 + 4147 +1011 + 16 +1012 DNA +1013 + Homo sapiens	
+320 + +323 + Hairpin ribozyme recognition site for PCNA	
<pre>-(400 \times 4147</pre>	1 ś
<pre>&lt;010 &gt; 4148 &lt;0211 &gt; 16 &lt;0210 &gt; DNA &lt;0213 &gt; Homo sapiens</pre>	
<220> <223> Hairpin ribozyme recognition site for PCNA	
<pre>&lt;4000 4148 aaattgtoad agadaa</pre>	16
<pre>&lt;010&gt; 4149 &lt;011&gt; 16 &lt;011&gt; DNA &lt;013&gt; Homo sapiens</pre>	
<pre><i220 -="" <i223="" for="" hairpin="" pcna<="" pre="" recognition="" ribozyme="" site=""></i220></pre>	
c4005 4149 titotgtoan caaatt	16
+.110+ 4150 +.111+ 16 +.112+ DNA +.113+ Homo sapiens	
-mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -mme- -me- -mme- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- -me- - -me- -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - -me- - - -	
-:400:- 4150 atotqqtota qttaac	16
<pre>#E100 4151 #E110 16 #E120 DNA #E120 Homo sapiens</pre>	

·(22) ·	
-0.023 - Hairpin ribozyme recognition site for PCNA	
<400 ± 4151	
thitigists thagaa	16
H210 + 4152	
H211 + 16	
HO10 MINA	
+1113 Homo sapiens	
+:2:20 +	
+CCC3 Hairpin ribozyme recognition site for PCNA	
<400 + 4152	
aaaqqgtot: gaotot	16
<210 % 4153	
<221 - 16	
KIDID KIND KIND KIND KIND KIND KIND KIND	
+C13+ Homo sapiens	
+:2.20>	
<pre><pre></pre> <pre>Hairpin ribozyme recognition site for lysyl oxidase</pre></pre>	
<:400 > 4153	
dagaagtaad tggtga	16
+:210 > 4154	
+211 × 16	
HOILS DNA	
+:213 · Homo sapiens	
\$20.0°	
<pre></pre> <pre>All?&gt; Hairpin ribozyme recognition site for lysyl oxidase</pre>	
+:400 + 4154	
otggagticae ogotgg	16
-01100- 4155	
+0110+ 16	
HIII. I I DNA	
-CLT30 Home sapiens	
+:00 0×	
Hairpin ribczyme recognition site for lysyl oxidase	
<400 > 415E	
eqeeegteae tggtte	16
₹21ar- 4156	

·:211 ·	l ô	
-:215 ·	DNA	
	Homo sapiens	
	·	
1230 A		
	Hairpin ribozyme recognition site for lysyl oxidase	
-(400 -	1156	
		1 ń
.4 - a.2 47	gibe ceagae	1.0
1 To 10 1 To	1167	
·(210 ·		
11 .		
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1.1.1.3	Homo sapiens	
43309 ×		
<0.033%	Hairpin ribozyme recognition site for lysyl oxidase	
<400>	41.57	
daggog	gtoca ogtacg	1 ń
-(2105	4158	
<211>		
<212>		
	Homo sapiens	
	T, Mo Oup 20110	
-:220×		
	Hairpin ribozyme recognition site for lysyl oxidase	
	Milipin likozymo rodognicion bico isi lybyl owradoc	
-(400 -	.1159	
		16
aaactq	gtotg godagt	エリ
<210 %	1160	
S2.0 / S211 +		
0012.		
* 1 3×*	Homo sapiens	
<pre></pre>		
HIII 2 31 +	Hairpin ribozyme recognition site for lysyl oxidase	
-(4000)		
titiatç	gtott gaagad	16
10100		
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1	DNA	
::::13:s	Hemo sapiens	
·:::::::::::::::::::::::::::::::::::::		
<225>	Hammerhead ribozyme recognition site for cdc 2 kinase	

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<pre>&lt;:400 &gt; 4160 tabaggtoaa gtggta</pre>	16
021:4:4161 0211:17 0211:5:DNA 0213: Homo sapiens	
$\cdot 22.0 \times \\ \cdot 22.3 \times \text{Hammerhead ribozyme recognition site for cdc 2 kinase}$	
-:400 - 4161 -aaatttotot attaaag	17
+010 + 4160 +011 + 16 +011 > DNA +013 > Homo sapiens	
<pre>&lt;220&gt; &lt;223 Hammerhead ribozyme recognition site for cdc 2 kinase</pre>	
+400 + 4162 agtoagtott caggat	16
<210> 4163 <211> 16 <212> TMA <213> Homo sapiens	
$\cdot$ 220% $\cdot$ 223% Hammerhead ribozyme recognition site for cdc 2 kinase	
<400 × 4163 ctggagtcag ctcgtt	16
H210 + 4164 H211 + 18 H212 - ENA H213 - Home sapiens	
shifts $+2.23 \times \text{Hammerhead ribozyme recognition site for cdc 2 kinase}$	
-400-4164 oqoqqaataa taagoogg	18
+0010+4165 +0010+18 +0010+DNA +0010+Bomo sapiens	

<pre>&lt;:225. &lt;:225 Hammerhead ribozyme recognition site for odd 2 kinase</pre>	
04 00 - 4165 ggaataataa googggat	18
<pre>0210 + 4166 0211 + 18 0213 + DNA 0213 + Homo sapiens</pre>	
Hammerhead ribezyme recognition site for odo 2 kinase	
H400 - 4166 geogggatot accatace	18
<pre>&lt;210&gt; 4167 &lt;211&gt; 18 &lt;212&gt; DNA &lt;213&gt; Home sapiens</pre>	
<pre>&lt;220&gt; &lt;223&gt; Hammerhead ribozyme recognition site for cdc 2 kinase</pre>	
<400 \ 4167 চল্লুপ্ৰtotas catascat	18
+0210	
+3220> +3223> Hammerhead ribozyme recognition site for cdc 2 kinase	
-04000-4168 totaccatac cattgact	18
SUL100-4169 -00118-18 -00100-DMA -00130-Hemo sapiens	
+:22:0:- +:2::3:- Hammerhead ribozyme recognition site for cdc 2 kinase	
+:400:4169	

-

cataccatty actaacta	18
+0010 + 4170 +0011 + 18 +0010 + DNA +0013 + Homb sapiens	
$\pm 220$ - $\pm 223$ - Hammerhead ribozyme recognition site for cdc 2 kinase	
-:400 - 4170 cattgastas ctatggaa	18
<pre> <pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre></pre>	
$<\!\!220^{\circ}\!\!\cdot\!\!$ $<\!\!223^{\circ}\!\!\cdot\!\!$ Hammerhead ribozyme recognition site for cdc 2 kinase	
<400 - 4171 yastaastat ggaagatt	18
<pre>&lt;210 &gt; 4172 &lt;2211 &gt; 18 &lt;2212 &gt; DNA &lt;2013 + Homo sapiens</pre>	
<pre>-unnos -unnos -unn</pre>	
<pre>&lt;400&gt; 4170 tqqaagatta taccaaaa</pre>	18
+C10> 4173 +C11> 18 +C12> DMA +C13> Homo sapiens	
*DESCRIPTION *DESC	
+44005 4175 qdaaqattat accaaaat	18
+0.100 4174 +0.110 18 +0.110 DNA +0.2102 Home samiens	

4220×		
-1.123 -	Hammerhead ribozyme recognition site for odc 2 kinase	
-(4.00)	1174	
		8
:U10 :		
-1211 -		
-1111-		
401134	Homo sapiens	
< (2.7.6) -		
	Hammerhead ribozyme recognition site for odc 2 kinase	
	•	
<4000 ≥	1175	
accaaa	atag agaaaatt 1	8
25155	117.5	
<210> <211>		
<212>		
	AMA Homo sapiens	
	TOMES SUPPLIES	
<220×		
k2230	ammerhead ribozyme recognition site for odo 2 kinase	
-:4005	176	
वृत्रवृत्रवत्रव	itty gagaaggt 1	8
-:210.s	177	
-11118		
·11.11.15		
	Homo sapiens	
. 1.2	Tombo Dupitono	
-:220:-		
H11135	ammerhead ribozyme recognition site for cdc 2 kinase	
	177	
-1400) 		0
ranaay	rtac ctatggag 1	0
-:[10:-	1:78	
-1111-		
.::12:-		
-00130-	lomo sapiens	
-000 00-		
·0.11 30*	lammerhead ribozyme recognition site for cdc 2 kinase	
-:400:-	178	
		19
	.cu. yyuyeeye	1 -
:2105	179	

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::211 · ::212 · ::213 ·		
H220 +	.  Hammerhead ribozyme recognition site for cdc 2 kir	nase
-(400 A tatqqa	4179 gagtty tgtataag	18
<pre>&lt;:210 + &lt;:211 + &lt;:212 + &lt;:213 +</pre>	. 18	
+:220 H +:223 H	Hammerhead ribozyme recognition site for cdc 2 ${ m kir}$	nase
-1400. agttgt	- 4180 gtgtat aagggtag	18
<pre>&lt;210 + &lt;211 + &lt;212 &gt; &lt;213 &gt; </pre>	· 18	
H220 N	· Hammerhead ribozyme recognition site for cdc 2 kir	nase
√4000 tigtat	· 418: gtataa gggtagac	18
<pre>si210 / si211 / si212 / si213 /</pre>	· 18	
-0.00 · -0.03 ·	· Hammerhead ribozyme recognition site for cdc 1. kir	ase
-:400.s ataaqg	s 418. Iggitad acacaaaa	18
+12101+ +12111+ +12121+ +12131+	- 18	
·0:20·		ıase

(400 × 4183 abasaabtad aggtbaag	18
<pre>4210 + 4184 4211 + 18 4212 + DNA 4213 + Homo sapiens</pre>	
নামার - নামার - Hammerhead ribozyme recognition site for odd 2 kinase	
<400 - 4184 otacagatca agtggtag	18
<pre>%210 + 4185 %211 + 18 %212</pre>	
+1220> +1223> Hammerhead ribozyme recognition site for odc 2 kinase	
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<400° 4186 aaaaaaatca gactagaa	18
<pre>%210% 4187 %211% 18 %212% DNA %212% Homo sapiens</pre>	
-CMMus- -CMMMs Hammerhead ribozyme recognition site for cdc 2 kinase	
-:400:- 4187 atcadactag aaagtgaa	18
+0010 + 4188 +0010 + 18 +0012 + DNA +0213 > Homo sapiens	

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4,22004
+223 · Hammerhead ribozyme recognition site for cdc 2 kinase
-1400 - 4188
ammagagtto ctagtact
                                                                       18
\pm 210 \pm 4189
-:211 - 18
HILL - DNA
+2213 + Homo sapiens
-100
+2.23 Hammerhead ribozyme recognition site for cdc 2 kinase
<400> 4189
aaygggttoo tagtactg
                                                                       18
<210> 4190
<211 > 18
<0.142 \times |DNA|
<213 \times Homo sapiens
<22.20 %
4223> Hammerhead ribozyme recognition site for cdc 2 kinase
H4005 4190
                                                                       18
gggttostag tactgcaa
\pm 210 \pm 4191
\leq 211 \leq 18
<212 > DNA
<1213> Home sapiens
-12.10 >
<22.3 Hammerhead ribozyme recognition site for cdc 2 kinase</p>
-:4000-4191
                                                                       18
ttoctagtac tgcaattc
H00100-4192
02110-18
H2121- DNA
+12.131 Homo sapiens
\pm 213: Hammerhead ribozyme recognition site for cdc 2 kinase
44000 4192
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actgca	attc	gggaaatt	18
·::11) >	4193		
-1.111			
-0.01.1.≯			
		sapiens	
-122119			
	Hamme	rhead ribozyme recognition site for cdc 2 kinase	
		ineda libobyme recognición bice isi eas xinabe	
<(400) >	1193		
		ggaaattt	18
10.0.1415.20		gguaree	10
si210 s	1191		
<0.11>			
<2112 >			
		sapiens	
10 a. J. J. C	כיוונטים	saprens	
<0.00>			
	1*	wheel wilders measurables site for ada 2 biross	
S.L.1.32	нашше	rhead ribozyme recognition site for cdc 2 kinase	
11000	1101		
<(40)0 >			1.0
ogggaa	latti	ototatta	18
4545	4100		
<210%			
<211>			
<212×			
42.133	Homo	sapiens	
-1220s			
<2235	Hamme	rhead ribozyme recognition site for cdc 2 kinase	
<400>			
वृवव्यव्यव्यव	ittta	totattaa	18
-11108			
42115			
42125			
412 1.515	$H \in m \ominus$	sapiens	
·121 U.S			
-122.31-	Hamme	rhead ribozyme recognition site for cdc 2 kinase	
-140úi.	4196		
qqaaat	ttct	ctattaaa	18
1.110	4197		
-C2115	15		
1	D1:A		
791 c.	Homo	saniens	

-: <u>22</u> 0 -	
$\pm 2.23 + \text{Hammerhead ribozyme recognition site for cdc } 2 \text{ kinase}$	
-:400 + 4197	
	18
aaatttotot attaaagg	10
H210> 4198	
+2211 + 18	
HOLD - DNA	
+2130 Homo sapiens	
K220 +	
$\pm 223 \pm \text{Hammerhead ribozyme recognition site for cdc } 2 \text{ kinase}$	
<400×4198	
	18
atttototat taaaggaa	10
<:210 ≥ 4199	
K211 N 18	
K212 DNA	
<pre>k213 Homo sapiens</pre>	
in the state of th	
<220%	
+3223 Hammerhead ribozyme recognition site for cdc 2 kinase	
,	
<400 + 4199	
ttototatta aaggaact	18
*:210.4 4200	
K2112-18	
+121218 DMA	
<213> Homo sapiens	
un section	
-CD230 Hammerhead ribozyme recognition site for odd 2 kinase	
+:400:+ 4000	
tototattaa aggaactt	18
s.ac.aa aggaasee	10
su 100 4001	
-0.11:-18	
HILLS DIA	
<pre>#ITIDE Homo sapiens</pre>	
The second secon	
+D100+	
+221.00 Hammerhead ribozyme recognition site for cdc 2 kinase	
+:4002 4201	
aaggaactte gteateea	18

·:210> 4202

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<111 - 19
HU12 - DNA
HDMD sapiens
-:2:1-
\pm 2.23 \times Hammerhead ribozyme recognition site for cdc 2 kinase
\pm 1400 \pm 4202
                                                                     18
aggaacting toathcaa
+0210 + 4203
\leq 211 + 18
HIIII - DNA
+213 · Homo sapiens
<1<u>220</u>→
<223> Hammerhead ribozyme recognition site for add 2 kinase
<400> 4203
aadttogtoa todaaata
                                                                     18
<210> 4204
<2117 18
RANTE > DNA
<213 > Homo sapiens
-1120-
-1.23 · Hammerhead ribozyme recognition site for cdc 1 kinase
\pm 400 - 4204
ttoutcator aaatatag
                                                                     18
+0.210 × 4.205
<211 > 18
+1212+ DNA
+213 > Homo sapiens
40.00
MARSS Hammerhead ribozyme recognition site for cdc 2 kinase
-:40C:- 4..05
                                                                     18
atccaaatat agtcagtc
+12100 41.06
000115-18
· ..1. · DNA
Homo sapiens
-112(H-
*2230 Hammerhead ribozyme recognition site for cdc 2 kinase
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03400 + 4206 Staaatslag toagtott	18
HD10 + 4207 HD11 + 18 HD12 + DNA HD13 + Homo sapiens	
$\pm 1.20 \gamma$ $\pm 1.23 \gamma$ Hammerhead ribozyme recognition site for cdc 2 kinase	
0400 + 4207 Batatagtoa gtottoag	18
<pre>0.100 4208 00112 18 00125 DNA 00135 Homo sapiens</pre>	
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<pre><pre><pre></pre> <pre><pre></pre> <pre><pre></pre> <pre></pre> <pre>Hammerhead ribozyme recognition site for cdc 2 kinase</pre></pre></pre></pre></pre>	
<pre>&lt;400 + 4209 gtcagtcttc aggatgtg</pre>	18
-0210 + 4016 -0211 * 15 -0210 * DNA -0.13 * Homo sapiens	
+00200+ +00200 Hammerhead ribozyme recognition site for cdc 2 kinase	
H400H 4010 idagictica ggatgtgc	18
<pre>*0:100+ 401! &lt;00110+ 16 &lt;02100+ DMA &lt;02130+ Homo sapiens</pre>	

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<220%
4223 Hammerhead ribozyme recognition site for cdc 2 kinase
-1400 - 4211
gatgtgctta tgcaggatt
                                                                     18
\pm 210 \pm 4212
\pm 2.11 \pm 13
AUC - DNA
-1213 · Homo sapiens
+12211+
+223 · Hammerhead ribozyme recognition site for odc 2 kinase
<400> 4212
                                                                     18
atgtgottat gcaggatto
<210> 4213
<211> 18
H212 > DNA
<213> Homo sapiens
41220×
<223> Hammerhead ribozyme recognition site for cdc 2 kinase
\pm 400 + 4013
                                                                    18
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<1.10 \pm 47.14
\pm 211 \pm 18
<21.1 DNA
<213 Homo sapiens
Hammerhead ribozyme recognition site for odc 2 kinase
-04000-4.114
                                                                    18
geaggatice aggittata
HI2100-4119
02110-18
-12111 DNA
+12130 Homo sapiens
-:22.30 Hammerhead ribozyme recognition site for cdc 2 kinase
-:400> 4Z15
```

ttocaggtta tatotoat	18
<pre>#210 * 4216 #211 + 18 #212 + DNA #213 + Homo sapiens</pre>	
+1220 + +1223 - Hammerhead ribozyme recognition site for cdc 2 kinase	
+:400 + 4216 topaggttat atotoato	18
+1210 × 4217 +1211 × 18 +2212 + DNA	
<213 * Homo sapiens	
<2238 Hammerhead ribozyme recognition site for cdc 2 kinase	
<pre>8400% 4217 daggttatat ctdatctt</pre>	18
<pre>&lt;010 * 4018 &lt;0011 * 18 &lt;0012 * DNA</pre>	
+2213 * Homo sapiens +220 *	
+223 $\times$ Hammerhead ribozyme recognition site for cdc 2 kinase +4400 $\times$ 4218	
qqttatatot catotttg +010-4019	18
<pre>&lt;011&gt; 16 &lt;012&gt; DNA &lt;013&gt; Homo sapiens</pre>	
+12.20:+	
$\pm 223\%$ Hammerhead ribozyme recognition site for cdc 2 kinase $\pm 400\%$ $\pm 421\%$	
thatatotoa totttgag <pre></pre>	18
+02110-18 +02120-DNA	
$\pm 213 imes$ Homo sapiens	

·220 ·	
4213 Hammerhead ribozyme recognition site for cdc 2 kinase	
+3450 × 4220	
tatotoatot tigagitt	18
<:210 * 4221 <:	
RE11 / 18 RE11 / DNA	
HANTA Homo sapiens	
* 1 * 11310 Sapiens	
K220 +	
+223 - Hammerhead ribozyme recognition site for cdc 2 kinase	
(400 < 4221)	
tetpatettt gagtttet	18
<210× 4222	
KZ11> 18	
<212> DNA	
+C13× Homo sapiens	
·220>	
<pre>v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;v&gt;</pre>	
s> namemented libozyme recognition site for edc 2 kinase	
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KD10 + 4223	
+D11 + 18	
SD1D - DNA	
+213+ Homo sapiens	
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HOLLOW DNA	
CLI30 Homo sapiens	
(IIII )	
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-(4)(i):- 4024	
tttgagtttc tttccatg	18
- Daylagence - eccepacy	10
221m2 1125	

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+213 - Homo sapiens
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\pm 2.25 \pm 1 Hammerhead ribozyme recognition site for cdc 2 kinase
+1400 > 4227
                                                                         18
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-01105 4008
<1111 18
RIZIO ENA
+2130 Home sapiens
+2223> Hammerhead ribozyme recognition site for cdc 2 kinase
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                                                                         18
ccaiqgaict gaagaaat
HD105 4/29
422115 18
\text{-}\text{CLITE} \text{-} \text{DMA}
·Mis- Home sapiens
+12201+
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+01105 4033 +0211+ 18 +0212+ DNA +0213+ Homo sapiens	
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gattistates etectggt
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Homo sapiens
(1) 3 (1) +
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intaticottoc tggtcagt
                                                                     18
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SIZIZY DMA	
SC1180 Homo sapiens	
scace	
+22239 Hammerhead ribozyme recognition site for cdc 2 kinase	
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+:210:+ 424:4	
+2212-18	
42120 DNA	
<pre><pre></pre> <pre></pre> <pre></pre> <pre>#213&gt; Homo sapiens</pre></pre>	

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+213 + Homo sapiens	
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001008 4049 00110-18 0010	
Home sapiens	
HARAGE Hammerhead ribozyme recognition site for cdc 2 kinase	
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27.2 m = 4.2 4 C	

+1210> 4246

923

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\pm 0.11 \cdot 13
HILL - DNA
HD13 - HDmb sapiens
Hammerhead ribozyme recognition site for cdc 2 kinase
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HILL FINA
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                                                                       18
agttatttat accaaatc
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1211 18
∹212 • DNA
+213 > Home sapiens
40000
40073 Hammerhead ribozyme recognition site for cdc 2 kinase
\pm 1400 + 4250
                                                                      18
itatitatad daaatoot
·210 · 4151
\leq 211 + 18
-1212 - ENA
+213 · Home sapiens
1.1
-00000 Hammerhead ribozyme recognition site for cdc 2 kinase
-14000 4.151
                                                                      18
caaatootac aggggatt
+.2100+ 4.52
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42230 Hammerhead ribozyme recognition site for cdc 2 kinase
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5210 5 32115		
4212 ×		
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KIJQUE. Kadal	Hammerhead ribozyme recognition site for cdc 2 kinase	
	Rammerhead Tibozyme recognition site for odd kinase	
-( <b>4</b> 00 -		
t. t gt at	tttq tcactcta	18
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·121.00-		
1.13	Homo sapiens	
<.22Ci+		
	Hammerhead ribozyme recognition site for cdc 2 kinase	
. 400.	W1.56	
		18
0.100		
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	Homo sapiens	
	To the second se	

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+1210 + 4258 +1211 + 18 +1212 + DNA +1213 * Homo sapiens	
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୍ୟମଣ୍ଡ 4259 agasgagtto ttpabaga	18
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+:210:- 4261 +:211:- 18 +:213:- Homo sapiens	
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+:400× 4261	

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H210m 4166 H211: 18 H212: DNA H213: Homo sapiens	

4220	Hammerhead	ribozyme	recognition	site	for	cdc	2	kinase	
	irananer ireaa	11202,1110	10009	0100		040			
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-12125									
	Homo sapien	S							
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	rammernead .	i ibozyme	recognition	sice	1 . I	Cac	-	KINASE	
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qqaacaa	atta aactggd	et							18
-1.11 01	4.170								
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	Homo sapiens	3							
*12200					_	,	_	, ,	
4121315 I	Hammerhead 1	ribozyme	recognition	site	ior	cdc	_	kinase	
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_									
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HL12 · DNA
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agatgatttt ggeettge
+1210 + 4272
\pm 211 + 18
· 21. • DNA
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<2200
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K4002 4272
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\le 2115 \cdot 18
KALIF DNA
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+1.7.7 () ×
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KC10 + 4278 KC11 + 18 KC12 + DNA KC13 + Home sapiens	
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-1400:- 4078 gaatacctat cagagtat	18
+01100-4179 +01110-18 +01100-DNA +00130-Hemo sapiens	
s(21.0) + s(22.0) Hammerhead ribozyme recognition site for cdc 2 kinase	
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HIII0 + 4281
4.211 \times 18
d211 → DNA
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44000 4284
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gaggtagtaa cactotgg	18
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HOLDS DNA	
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+:220 s	
$\pm 223 imes$ Hammerhead ribozyme recognition site for odo 2 kinase	
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cagotogita otoaacto	10
-CD1CO41-93	
+211> 18	
HOLDS DIA	
82133 Homo sapiens	
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·(2.30)·	
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agetegttae teaaetee	18
H210:4294	

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                                                                        18
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·(220) -
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·III1.1 > DNA
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H2205
<123> Hammerhead ribozyme recognition site for cdc 2 kinase
+1400 > 4505
                                                                        18
taqcaactaa gaaaccat
+2210> 430€
\pm 0.115 \cdot 18
<212> DNA
<!li>113> Homo sapiens
\pm 2135 Hammerhead ribozyme recognition site for cdc 1 kinase
-:400:- 4306
                                                                        18
qaaaccattt tccatggg
-02100-4307
022110 18
+02123 DNA
+22130 Homo sapiens
+2130 Hammerhead ribozyme recognition site for cdc 2 kinase
-:400> 4:07
```

aaaccatttt coatgggg	18
+1310 + 4308	
×:211 + 18	
H212 + DNA	
H213 Homo sapiens	
- Live Home bupicino	
+(220 ·	
+2223 Hammerhead ribozyme recognition site for cdc 2 kinase	
name name industries in the control of the control	
+(400 × 4308	
aaccattito catgggga	18
aassace tes caegggga	10
±210 ± 4309	
×211 + 18	
<212 - DNA	
<213 - Homo sapiens	
CLIP TOMES Suprems	
*13.70 -	
Hammerhead ribozyme recognition site for cdc 2 kinase	
".LLD" hammerhead fibozyme recognition site for cdc 2 kindse	
×(400 + 4309	
	18
accattttcc atggggat	10
1010. 1010	
+:210 × 4310	
+1211 × 18	
HIM KIMA	
<pre><pre></pre> <pre>#INT3 &gt; Home sapiens</pre></pre>	
+10.20 %	
${\it clids}$ Hammerhead ribozyme recognition site for cdc 2 kinase	
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atggggatto agaaattg	18
*:210 · 4:311	
+1711 + 18	
KIND DIA	
<pre>#2175 Homo sapiens</pre>	
<220.*	
+22.86 Hammerhead ribozyme recognition site for cdc 2 kinase	
+:40005-4511	
tqqqqattca gaaattga	18
+02100 4312	
+00110+ 18	
4:212: DNA	
<213> Homo sapiens	

+:2200+	
+:400 + 4312 agattgatoa actottoa	18
+010 + 4313 +011 + 18	
HULL DNA	
HU13 - Homo sapiens	
+1220 +	
$\pm 223 \pm { m Hammerhead}$ ribozyme recognition site for cdc 2 kinase	
+(400× 4313	
gatdaactot toaggatt	18
+210× 4314	
×2213 18	
<pre><pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre><pre></pre>&lt;</pre></pre>	
+213> Homo sapiens	
42209	
$<\!22238$ Hammerhead ribozyme recognition site for add 2 kinase	
×400> 4314	
toaactetto aggatttt	18
<210 > 4315	
4211× 18	
%212 % FNA	
<213 > Homo sapiens	
KD20 +	
$\pm 0003$ · Hammerhead ribozyme recognition site for odd $\Omega$ kinase	
+:400 × 4315	
ticaggattt toagaget	18
·(210 + 4316	
(211. 18)	
-02120 DNA	
-:213: Homo sapiens	
+(2/2 0)+	
+2233+ Hammerhead ribozyme recognition site for cdc 2 kinase	
·(400) · 4316	
teaggattitt cagagett	18
2010: A017	

H210H 4317

```
4211 - 13
-.212 - DNA
-213 · Homo sapiens
-(220) +
\pm 223 \pm \text{Hammerhead ribozyme recognition site for cdc } 2 \text{ kinase}
+(400 + 4317)
                                                                          18
daggattite agagettt
<210 → 4318
\pm 2119 \cdot 18
4010 - DNA
\cdot 213 \cdot Homo sapiens
<220>
<223> Hammerhead ribozyme recognition site for cdc 2 kinase
<400> 4318
aggattttca gagetttg
                                                                          18
<.310 > 4319
<211> 16
<212> DNA
<213 > Homo sapiens
<10.00 ×
+2223 Hammerhead ribozyme recognition site for cdc 2 kinase
-:400 · 4319
thagagettt gggeacte
                                                                         18
\pm 210 \times 4320
\pm 1.711 \pm 18
471.1 \times DNA
+1113 > Homo sapiens
+1220×
-:223 Hammerhead ribozyme recognition site for cdc 2 kinase
+:400 + 4320
dagadetitg ggdadted
                                                                         18
<2100 4321
401111-18
-1.1. DNA
*Clime Hume sapiens
·17.200 ·
<2230 Hammerhead ribozyme recognition site for cdc 2 kinase</p>
```

-

<pre></pre>	18
<pre>4310 + 4322 4311 + 18 4312 + DNA 4313 + Homo sapiens</pre>	
<pre><pre></pre> <pre>&lt;223 * Hammerhead ribozyme recognition site for cdc 2 kinase</pre></pre>	
-(400) + 4322 -stoosaataa tgaagtgt	18
<pre>&lt;210 + 4323 &lt;211 * 18 &lt;212 * INA &lt;213 * Home sapiens</pre>	
K220.	
<223 Mammerhead ribozyme recognition site for cdc 2 kinase <400> 43.33 agtugaatet ttacagga	18
+210> 4324 +211 + 18 +211 > DNA +213 + Home sapiens	
<pre>&lt;2000 s &lt;2003 Hammerhead ribozyme recognition site for cdc 0 kinase</pre>	
<pre>04005 4324 tggaatcttt acaggact</pre>	18
+2010:- 4325 +2212: 18 +2212: DNA +2213: Homo sapiens	
<pre>&lt;2208 &lt;2235 Hammerhead ribozyme recognition site for cdc 2 kinase</pre>	
<4005 4325 qqaatottta daggadta	18
+0.100+432t +0.110+15 +0.120+DNA +0.2130+Homo sapiens	

```
+(2]:) +
+223 - Hammerhead ribbzyme recognition site for odd 2 kinase
<400 + 4326
gaatetttae aggaetat
                                                                       18
+2210 + 4327
\pm 311 \pm 18
+121.1 + DNA
+213 · Homo sapiens
k(220) k
4223. Hammerhead ribozyme recognition site for cdc 2 kinase
4400 → 4327
                                                                      18
acaggactat aagaatac
<210 > 4328
<2115 18
4212 > DNA
<:213> Homo sapiens
<(220 s
<223 > Hammerhead ribozyme recognition site for cdc 2 kinase
+3400 \times 4328
                                                                      18
aqqactataa gaatacat
+1210 × 4329
-111> 18
<010> DNA
<113> Homo sapiens
41210.4
\pm 2238 Hammerhead ribozyme recognition site for cdc 2 kinase
<4000 4329</p>
                                                                      18
ataaqaatad atttccca
*IN10: 4330
·11111 18
<math DNA</pre>
+0.0130 Homo sapiens
\pm 21.30 Hammerhead ribozyme recognition site for cdc 2 kinase
-:400> 4330
```

gaatacattt eccaaatg	18
<pre><pre></pre></pre>	
-:210 + 4331	
+(211 + 1d	
+1313 + DNA	
+213× Homo sapiens	
-1220 -	
-MANGE Hammerhead ribozyme recognition site for cdc 2 kinase	
+(400 > 4331	
aatacatttc ccaaatgg	18
+1210 + 4332	
<211 - 18	
<212> DNA	
<pre>&lt;013&gt; Homo sapiens</pre>	
K220 S	
<223> Hammerhead riboxyme recognition site for cdc 2 kinase	
<ul><li>400 × 4332</li></ul>	
atadatttod daaatgga	18
+210 × 4333	
+211> 18	
+1212 - DNA	
+013 + Homo sapiens	
+12200 ×	
$\pm 2.2.38$ Hammerhead ribozyme recognition site for cdc 2 kinase	
K4005 4333	
ggaagootag catoocat	18
ggangootag oddoosad	
H2108 4334	
x0110 18	
HOMOS DNA	
<pre>&lt;213&gt; Home sapiens</pre>	
•	
+:22(c)	
$\pm 2238$ Hammerhead ribozyme recognition site for cdc 2 kinase	
+:400:> 4334	
octagoatoc catgtoaa	18
(110) 11 25	
+02100+ 4035	
-0.110- 18	
+Ch12:+ DMA	
Homo sapiens	

<pre>::220 + ::::::::::::::::::::::::::::::::::::</pre>	
(400 - 4335 tessatytoa aaaasttg	18
KL10 + 4336 KL11 + 18 KL12 + DNA	
·213 · Homo sapiens	
<2200 · <223 · Hammerhead ribozyme recognition site for odd 2 kinase	
୍ୟପତ୍ତ 4336 Caaaaacttg gatgaaaa	18
<210 - 4337 <211 - 18 <212 DNA	
<pre>kill3 + Homo sapiens</pre>	
<pre>%ABO + @BBS + Hammerhead ribbzyme recognition site for cdc 2 kinase</pre>	
0400 = 4337 asatqqcttq gatttqct	18
0.110 × 4338 0.111 × 18 0.117 → DNA	
2213 * Homo sapiens	
<pre>&lt;2200 * .223 * Hammerhead ribozyme recognition site for cdc 2 kinase   </pre>	
0400 × 4338 gottagattt gototoga	18
1210 - 4039 1211 - 18 1212 - DNA	
11130 Homo sapiens	
MING Hammerhead ribozyme recognition site for cdc 2 kinase	
(40):- 4539 Stiggatity ototogaa	18
721d1. J14d	

```
-:211 · 13
-1112 · DNA
·113 · Homo sapiens
+12.20 ×
\pm 1235 Hammerhead ribozyme recognition site for cdc 2 kinase
-:400× 4340
                                                                       18
gatftgctct cgaaaatg
+0210 \times 4341
-211 - 13
4212 + \text{DNA}
+213 > Homo sapiens
x220 ×
<223 Hammerhead ribozyme recognition site for cdc 2 kinase</p>
<4005 4341
tttgctctcg aaaatgtt
                                                                       18
+:210:- 4342
\pm 211 \times 18
+212> DNA
+:213> Home sapiens
<1220 >
+1223> Hammerhead ribozyme recognition site for cdc !! kinase
3400> 434D
qaaaatgtta atctatga
                                                                       18
<210 - 4343
<2211 · 18
<212 - DNA
+:213 · Homo sapiens
· (1 -
-CDB - Hammerhead ribozyme recognition site for cdc D kinase
-1400 - 4343
aaaatgttaa totatgat
                                                                      18
HI2100 4344
H2115 18
HILLI DNA
+2213 - Home sapiens
-:2201-
<223 Hammerhead ribozyme recognition site for cdc Z kinase</p>
```

943

<pre>::400 + 4344 styttaatot atgateca</pre>	18
<pre>#210 * 4345 #211 * 18 #2212 * DNA #2213 * Homo sapiens</pre>	
$<\!\!2208$ $<\!\!2208$ Hammerhead ribozyme recognition site for odd 2 kinase	
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<pre><c10 +="" 4346="" <c11=""> 18 <c1c <c13="" dna=""> Homo sapiens</c1c></c10></pre>	
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<pre></pre> <pre>&lt;4000 4347 aaacqaattt ctggcaaa</pre>	18
+02100 4348 +02110 18 +02120 DNA +02130 Homo sapiens	
+2200+ +2230 Hammerhead ribozyme recognition site for cdc 2 kinase	
-:400:- 434- aacqaattto tggcaaaa	18
+0.100+4+49 +0.110+18 +0.110+ DMA +0.215> Homo sapiens	

$\cdot \cdot \cdot$ Hammerhead ribbzyme recognition site for cdc 2 kinase	
-1400 + 4349 abyaatttot ggcaaaat	18
+1210 + 4350 +1211 + 18 +1212 + DNA +1213 + Homo sapiens	
$\pm 220$ - $\pm 223$ - Hammerhead ribozyme recognition site for cdc 2 kinase	
<pre>&lt;:400 + 4350 cactgaatca tocatatt</pre>	18
<pre>#2108 4351 #2118 18 #2128 DNA #213 Homo sapiens</pre>	
<2220> <2235 Hammerhead ribozyme recognition site for cdc ( kinase	
<400 × 4351 tquatcates atatttta	18
<pre>&lt;010 + 4352 &lt;011 + 18 &lt;0110 * DNA &lt;2113 * Homo sapiens</pre>	
R220 R223 Hammerhead ribozyme recognition site for cdc 1 kinase	
k:400% 4352 teatecatat titaatga	18
+0010% 4353 +0010% 18 +0010% DNA +0013% Homo sapiens	
+2200> +223> Hammerhead ribozyme recognition site for cdc 2 kinase	
+:400> 4353	

atopatattt taatgatt	18
<pre>%210 * 4354 %211 * 18 %212 * DNA %213 * Homo sapiens</pre>	
<pre><pre>&lt;0.020 + &lt;0.023 - Hammerhead ribozyme recognition site for cdc 2 kinase</pre></pre>	
0400 - 4354 topatatttt aatgattt	18
H210 + 4355 H211 + 18 H212 > DNA H213 - Homo sapiens	
<pre>&lt;220 </pre> <pre>&lt;223 Hammerhead ribozyme recognition site for cdc 2 kinase</pre>	
<pre></pre> <pre>&lt;400 · 4355</pre> <pre>coatatitta atgatttg</pre>	18
<pre>&lt;210 + 4356 &lt;2211 + 18 &lt;2112 + DNA &lt;2013 + Homo sapiens</pre>	
<pre><d200> <d2038 2="" cdc="" for="" hammerhead="" kinase<="" pre="" recognition="" ribozyme="" site=""></d2038></d200></pre>	
<400> 4356 catattttaa tgatttgg	18
+02105 4357 +0.110+ 18 +02100+ DNA +02130+ Homo sapiens	
+:220:- +:2235 Hammerhead ribozyme recognition site for cdc 2 kinase	
+44005 4357 thaatgaitt ggacaatc	18
+0:10: 4:58 +0:11: 18 +0:12: DNA +0:13: Homo sapiens	

+(225 × +(235 ×	Hamme	erhead	ribozyme	recognition	site	for	ede :	? kinase	
(400) taataa		gacaat	Ca						18
		gacaac	.ca						10
+1210 + 4211>									
<212 × <213 ⋅		sapien	ıs						
-1220 %									
+12233 +	Hamme	erhead	ribozyme	recognition	site	for	ede :	2 kinase	
<400×									1.0
togaca	aatca	gattaa	ıga						18
<0.10>									
<21125 <21125									
		sapien	ıS						
₹220÷									
KI223.F	Hamme	rhead	ribozyme	recognition	site	for	ede 2	2 kinase	
<400%									
gaagat	gtag	ctttct	ga						18
-210 -									
<211 · < < 212 · <									
		sapien	S						
<2200 s									
-1723 ·	Hanme	rhead	ribozyme	recognition	site	for	cycli	in B1	
-(400)	4361								
toogaç	gticac	caggaa							16
-2100-									
· 1 111 · · 21.11 ·									
		sapien	S						
·1702.00									
	Hamme	rhead	ribozyme	recognition	site	for	cycli	in B1	
·(4 )-01-	4062								
ccaq*.ç	jtota	agccag							16
.:21e.	1363								

<pre>&lt;211 * &lt;2.12 * &lt;1.13 *</pre>	DUIA	sapier	ıs					
-1220 + -1223 +	Hamme	erhead	ribozyme	recognition	site	for	cyclin	B1
(400 ) catqt		gctttc	2					1.
<pre>&lt;:010 + &lt;:211 +</pre>								
-:212 + -:213 +		sapier	ìs					
<200 + <223 +	Hamme	erhead	ribozyme	recognition	site	for	cyclin	B1
<400. aagcag		accaaa	ì					1.
<210×	16							
40124 42134		sapier	ıs					
40006 40035	Hamme	rhead	ribozyme	recognition	site	for	cyclin	B1
c400 s actggg		gaagto						1 (
<pre>%210&gt; %211&gt; %211&gt;</pre>	16							
-12238		sapier	ıs					
		rhead	ribozyme	recognition	site	for	cyclin	B1
-:4(mi∷- tdactç		cattat	:					1 (
·:::10:- ·:::11:- ·:::1:::	1 €							
:01:05 :02:05	Homo	sapien	15					
	Hamme	rhead	ribozyme	recognition	site	for	PCNA	

·:400 ·	4367	
वर्ग्य	gtima gggoto	16
-:210 ·	4368	
-:211		
-:212 ·		
	Homo sapiens	
	HOMO Sapiens	
·::::20 -		
-1.2.2.3 -	Hammerhead ribozyme recognition site for PCNA	
(400 ·	4368	
	gtodo adgiei	16
.i	good dogodo	
·1210>		
<211>		
<212 -		
<213>	Homo sapiens	
<2200 ×		
	Hammerhead ribozyme recognition site for PCNA	
<4005	4369	
atqaqq	gtictg aggget	1 6
<1.U>	4370	
<.111	16	
G2125	DNA	
	Hemo sapiens	
0220%		
<2223>	Eammerhead ribozyme recognition site for PCNA	
-14005 -14005	1276	
		16
	gthad caaatt	10
-1175-	4371	
-1111	1 rî	
	DNA	
	Homo sapiens	
-000 00-		
-11 1130	Hammerhead ribozyme recognition site for lysyl oxidase	
-14 û () ()	4.71	
	stace tggtge	16
	4.22	
100		
ahli:		
1.1.		
:21 <i>5</i> >-	Homo sapiens	

-0220 · -0323 ·	Hammerhead	ribozyme	recognition	site	for	lysyl	oxidase	
-(400 - ctgga	4372 gtoad ogotgo	3						16
<pre>%210 + %211 + %212 + %213 +</pre>	1 กั	ıs						
-:220 - -:223 -	Hammerhead	ribozyme	recognition	site	for	lysyl	oxidase	
<400× ogoco	4373 gtcac tggttd	2						16
<210 + <211 + <212 + <213 +	16	1S						
<220% <2235	Hammerhead	ribozyme	recognition	site	for	lysyl	oxidase	
<400> gtacgo	4374 gtoto deagad	:						16
<pre>(0.10% &lt;0.11% &lt;0.1</pre>	16	ıs						
:2200 :2236	Hammerhead	ribozyme	recognition	site	for	lysyl	oxidase	
-:4005 ⊝aggog	4375 gtoda ogtacç	1						16
-::105 -::11:- -::2125 -::2135	1 r.	ıs						
-12200- -12230-	Hammerhead	ribozyme	recognition	site	for	lysyl	oxidase	
:400>	4.576							

aaastgtotg gooagt	16
<pre>1210 + 4377 1211 + 16 1212 + DNA 1213 + Homo sapiens</pre>	
$\pm 2200 + \pm 223$ . Hammerhead ribozyme recognition site for lysyl oxidase	
<pre><idu0> 4377 tttotgtott gaagac</idu0></pre>	16
<pre>KD10 + 4378</pre>	
<pre>&lt;220 * &lt;223 * Representative Haipin Ribozyme</pre>	
<pre>6400. 4378 aacqagctag aaccagacca gagaaacaca cgttgtggta tattacctgg ta</pre>	52
K2108 4379 K2114 55 K2129 DNA K213 - Homo sapiens	
<pre><!--220% <!223 - Fepresentative Haipin Ribozyme</pre--></pre>	
<pre>&lt;:400&gt; 4379 ctggctcaag aactggacca gagaaacaca cgttgtggta tattacctgg ta</pre>	52
H2108 4380 H2118 50 H2125 DNA H2135 Homo sapiens	
+00000+ +0000> Representative Haipin Ribozyme	
-(400)- 4080 ocaqogqtaq aaccagacca gagaaacaca cgttgtggta tattacctgg ta	52
+0210:- 4081 +0210:- 51 +0210:- DMA +0213> Homo sapiens	

```
-:22u→
<!223 Representative Haipin Ribozyme</pre>
+1400 + 4381
                                                                     52
ageoctoaag aageagacca gagaaacaca egttgtggta tattacetgg ta
\pm 210 + 4382
<211 - 40
-1211 - DNA
All: Artificial Sequence
4.1.10 ×
HALLS + Synthetic human hammerhead ribozymes targeting PCNA
<\!400+4382
                                                                 40
gagecetgeu gaugageaat tttttgegaa aaccaggege
<210> 4383
<211≥ 38
KANAN DNA
H220×
+223> Synthetic human hammerhead ribozymes targeting PCNA
+3400> 4383
                                                                     38
agoccugoug augaggoogt aaggoogaaa ocaggogo
+210 > 4384
\pm 211 \leq 38
-1212 - DNA
<213 Artificial Sequence</pre>
<11.20 ×
<223 - Synthetic human hammerhead ribozymes targeting PCNA</p>
+14000 4384
                                                                     38
agoccurcug augaggoogt aaggoogaaa ocaggogo
HITTOH 4385
32111-39
·IIII DNA
*Ill: Homo sapien
+07.53+ General structure of chimeric DNA/RNA ribozyme
\{(x,y)\in X_{k+1}(x)\}
·IIII: modified base
-11111 (1..8)
%223> Where n is a, c, g or t
```

.

```
40000 ×
MACL * modified_base
MACL * (33..39)
-0.23 Where n is a, c, g or t
-1400 + 4385
                                                                              39
nnnnnnncu gaugagcaat tttttgcgaa aannnnnn
\pm 1210 \pm 4386
4211 · 16
HILL - DNA
+1113 + Homo sapiens
+0.5790 \times
....3 · General target RNA of chimeric DNA/RNA ribozyme
<1220 >
<221> modified base
4222 > (1..8)
<223> Where n is a, c, g or u
<2220 ×
<221 - modified base
<2222 (11..16)
H223 Where n is a, c, g or u
74005 4386
                                                                        16
nnnnnnnnhu nnnnn
H2105 4387
-1211> 51
+1212> DNA
+213> Homo sapiens
<!!!!
() >
+00.33 Representative hairpin ribozyme
ST11014
+DDD: modified_base
<DDD: (1..8)</pre>
\times 1230 \cdot Where n is a, c, g, t or u
-:2200-
-:221: modified base
<22225 (13..16)</pre>
\cdot \text{2230} \cdot \text{Where n is a, c, g, t or u}
-14001- 4:37
                                                                              52
nnnnnnnnag aannnnacca gagaaacaca cguuguggua uauuaccugg ua
12:10: 4:88
<211> 16
```

```
HO12: DNA
-:213 · H mo sapiens
·(226 ·
+223 · Target RNA of representative hairpin ribozyme
-m20 -
-m21 - modified_base
-m220 - (1..3)
-m23 - Where n is a, c, g, or u
·[220 ·
+221 \times \texttt{modified\_base}
:1222 · (5:
·223 · Where n is a, c, g, or u
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nnnsnguenn nnnnnn
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        -1220 -
        -:223 · MMP-1 ribozyme recognition site
        <400 + 4389
 gtggtgtoto acaget
                                                                                              16
        K2105 4390
        +1.111 - 16
        ·IIII DNA
        -17130 Homo sapien
        \star (\underline{\mathbb{Z}} \, \underline{\mathbb{Z}} \, \Omega) \star
        +:2130 MMP-1 ribozyme recognition site
        -04000-4390
                                                                                              16
 agtitighedt captga
        HIL100-4391
        -22115 16
-22125 DNA
        400130 Homo sapien
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        <4000 4591
```

ccaaggtoto tgaggg	16
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SUL102 4394 SUL103 16 SUL103 DNA SUL133 Home sapien	
<pre><coor< pre=""> <pre><coor< pre=""> <pre><coor< pre=""> <pre>MMP-1 ribozyme recognition site</pre></coor<></pre></coor<></pre></coor<></pre>	
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-:400+ 439+ gaactgtora ttttot	16
+1210 + 439* +1211 > 16 +1212 + DNA +1213 + Home sapien	
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-:400:- 439r totoagtoat tittaa	16

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+0010 = 4395 +0010 + 16 +0010 + DNA +0013 + Homo sapien	
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KD10F 4402	

4212 ← DNA 4213 ← Homo	sapien			
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<pre><dub -="" 4402="" gtotggtbba="" pre="" tatgaa<=""></dub></pre>	a			16
+1010 + 4403 +1011 + 16 +1112 + DNA				
*1213 + Homb *1220 + *1223 + IL-1	beta ribozyme	recognition	site	
<400 - 4403 tocatgreet tigtac				16
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+1210+ 4405 +1211+ 16 +1212+ DNA +12131 Homo	sarien			
+2205 +2235 IL-1	beta ribozyme	recognition	site	
4000- 4405 gatttgtott baabaa	ì			16
+:210:5 4406 +:011:5 16 +:011:5 DMA +:013:5 Hemo	sarien			
+:2:00:5 +:2:23:- II:-1	beta ribozyme	recognition	site	
<pre>&lt;:400:- 4406 tttgagtctq cchagt</pre>	:			16
*:210:+ 4407 *:211:- 1* *:212:- DNA *:213>- Homo	sapien			

+:220 + +:223 + IL−	-l beta ribozyme recognition site	
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+0010 + 440 +0010 + 16 +0010 + DNA +0110 + Hom	A	
::226.÷	-l beta ribozyme recognition site	
<pre>+:400 + 440 tttgtgtstt ccta</pre>	)÷	16
<pre><di0> 440 <di1> 16 <di1> 5NA <d13> Hom</d13></di1></di1></di0></pre>	) <del>,</del>	
-(220%	-l beta ribozyme recognition site	
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+(220)+ +(223)+ IL+	-1 beta ribozyme recognition site	
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+0010 + 441 +0011 + 16 +0010 + DNA +0013 + Hom	A	
+:120.+ +:125.+ VEG	GF ribozyme recognition site	
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+0:108 441 +0:110 16 +0:100 DNA +0:150 Hom	A	
+:220×		

	-: 2.2.3 > VEGF ribozyme recognition site	
	(40) + 4412 topo aggotg	1 6
	8210 × 4413 8211 × 16	
	RANG + DNA	
	<pre>%213 Homo sapien</pre>	
	(200 )	
	HILIBA VEGF ribezyme recognition site	
	(400 × 4413	
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	(£10) 4414	
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	<pre></pre> <pre>&lt;</pre>	
	6400 - 4414  otgt ottgoattgo a  1. The state of the s	21
	\$ 210 % 4415	
	\$210, 4415 \$211, 21	
	ROLLS DNA	
	<pre><!--!document</td--><td></td></pre>	
	(220)	
	<pre>&lt;223 * IL2 ribozyme recognition site</pre>	
	0400> 4415	
ttgcac	ttgt cacaaacagt g	21
	स्प्र10% 4416	
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	C.130 Homo sapien	
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	CLLS: IL2 ribozyme recognition site	
	::400:- 4416	
	grigit otagaagaag a	21
	02108 4417	
	C2118-21	
,	C2128 DNA C2130 Homo sapien	
	(220)	
	223> IL2 ribczyme recognition site	

्य )। र 4417 taccttrtgt casagcatca t	21
0210 + 4418 0211 + 21 0212 + DNA 0213 + Homo sapien	
-1226 1NF-gamma ribozyme recognition site	
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+:210 + 4419 +:211 + 21 +:212 + DNA +:213 + Homo sapien	
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<pre>&lt;0.00 + &lt;0.03 + INF-gamma ribozyme recognition site</pre>	
k400 - 4420 aaaaagqaqt caqatgetgt t	21
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02008 00088 INF-gamma ribozyme recognition site	
-:40(8 4421 gtaatggttg tootgeetge aa	22
+00100 4407 +00110 01 +00125 DNA +00135 Homo sapien	
-:220 -:223- INF-gamma ribozyme recognition site	
<pre>&lt;:400&gt; 4422 ctgtgactgt ctcacttaat c</pre>	21

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HIII) - 4423
       HL11 - 21
       \pm 1.111 \pm \text{DNA}
       -1113 · Homo sapien
       -:225 -
       ::12: INF-gamma ribozyme recognition site
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                                                                                     21
ttgaatqtqt caqqtgaccc t
       \leq 2.19 \times 4424
       \cdot 1211 \cdot 21
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                                                                                     .21
gogtttgagt cagcaaagaa g
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       +1010 + 4406
       \times 2.111 + 2.11
       SINIE DNA
       +22135 Homo sapien
       +2230- IL-1 alpha ribozyme recognition site
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      +0110+4407
+011+10+11
+0110+DNA
       +00130 Homo sarien
       00000 IL-1 alpha ribozyme recognition site
       -(400)- 4427
ggtctggag* ctcacttgtc t
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0011 + 01 0010 + DNA 0013 + Homo	o sapien	
<1220 + <1223 + IL-1	l alpha ribozyme recognition site	
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0210 + 4429 0211 + 21 0212 + DNA 0213 * Homo		
+:220× +:223× IL-1	alpna ribozyme recognition site	
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+:0200+ +:023+ IL-1	alpha ribezyme recognition site	
+:4005 4431 agtaattqqt occdat		21
-0.160-4430 -00110-01 -01100-DNA -02130-Homo		
+:210:+ +:213:+ 1L-1	alpha ribozyme recognition site	
-(400)- 443) cogtgotggt otoga-		21
+0:10:+ 4433 +0:11:+ 21 +0:12> DNA		

	GC13 + Homo sapien	
	K000 +	
	- CLBB - IL-1 alpha ribozyme recognition site	
	(400 + 4433	2.1
aatc	ctdagt cagoogtgtt t	21
	<210 + 4434	
	-0.11 + 18	
	HO12 - DNA	
	+213 * Homo sapien	
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	3223 · IL-1 alpha ribozyme recognition site	
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	eg 1555 dage gada	
	<210 * 4435	
	<211 > 21	
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	<pre>&lt;223&gt; IL6 ribezyme recognition site</pre>	
	<4005 4435	
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	H210.5 4436	
	82118 21 	
	-2125 DNA <2135 Homo sarien	
	215 Homo Suprem	
	NULPON TO THE PROPERTY OF THE	
	<pre>&lt;2223&gt; IL6 ribezyme recognition site</pre>	
	<4008 4436	
catca	actogt ctitiggagt t	21
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	SCIOS DNA	
	-00135 Homo sapien	
	<pre>&gt;C265 &gt;C235 IL6 ribezyme recognition site</pre>	
	Z The liberyme recognition site	
	40054437	
ttaat	tgdagt ddagddtgag g	21
	H2105-4438	
	-02118 A1	
	-021105 DNA	
	-213. Homo sanien	

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<pre>&lt;:22↑ * &lt;:dd &gt; 1L6 ribbzym</pre>	e recognition :	site	
(400 + 4438 ttottotyyt pagaaacoty t			2:
H218 + 4439 H211 + 21 H212 + DNA H213 + H0mb sapien			
<0.00 ≠ <0.03 + 116 ribozyme	e recognition s	site	
-4400 - 4439 agaaacstyt csastgggda c			21
<pre></pre>			
<220 -333 - IL8 ribozyme	e recognition s	site	
+400+ 4440 ttgtgtaggt ctgttgtagg g			21
+210 + 4441 +211 > 21 +212 + ENA +213 > Homo sapien			
∘020° ∘000° IL8 ribosyme	recognition s	site	
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<pre>\$210 &gt; 4442 \$211 &gt; 21 \$212 &gt; DWA \$213 &gt; Homo sapien</pre>			
400C ⋅ +0000 IL8 ribozyme	recognition s	;ite	
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0010 + 4443 0011 + 21 0012 + DNA 0013 + Homo sapien			
<220 · K223 · MMP-2 ribozy	me recognition	. site	

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H400+ 4443 googogoagt ogascatoat o	21
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<pre>&lt;(220 * &lt;0.03 * MMP-2 ribozyme recognition site</pre>	
<pre>&lt;400 + 4444 godaagt pgt cogtgtgaag t</pre>	21
<pre>%210 + 4445 %211 + 21 %212 &gt; DNA %213 + Homo sapien</pre>	
<pre>&lt;220 &gt; &lt;223 &gt; MMP-2 ribozyme recognition site</pre>	
-400-4445 oggottetgt occcatgaag o	21
+010 × 4446 +011 × 01 +010 × DNA +013 × Homo sapien	
<pre>+0.00 * +0.03 * MMP-D ribozyme recognition site</pre>	
H4005 4446 dedectgtgt ottobootto a	21
+(210> 4447 +(211> 21 +(212> DNA +(213> Homo sapien	
HIMBS MMP-0 ribozyme recognition site	
(400% 444°) gaacttoogt otgtoocagg a	21
02105 4448 02115 21 02115 DNA 02135 Homo sapien	
K22C> K22C> MMP-2 ribozyme recognition site	
€400≥ 4448	

ttccgtstyt ccsaqyatga c	.21
0210 + 4440 0211 + 21 0212 + 5MA 0213 + Homo sapien	
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<pre>kD16: 4453 cD11: D1 cD12: DMA cD13: Heme sapien</pre>	
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(400): 445: ttotttgggt ottgttttt t	21

	0210 + 4454 0211 + 20 0212 + DNA 0213 + Homo sapien	
	K120 · K123 · MMP-3 ribozyme recognition site	
	<pre>&lt;400 + 4454 adub ttpcaatoot</pre>	.20
	<pre>&lt;210+ 4455 &lt;211+ 21 &lt;212+ DNA &lt;213+ Homo sapien</pre>	
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	ollos Clas MMP-3 ribozyme recognition site	
	:400> 4457 gags etetiteact e	21
	12105 4455 12115 21 12125 ENA 12135 Homo sapien	
	(220) (213) MMP-3 ribozyme recognition site	
	:400% 445% stgt otdaagatga t	21
	0.108-4459 (211)-21	

	COLOR DNA	
	<pre>&lt;0.13 · Homo sapien</pre>	
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	d2.2.0 →	
	<pre><dd3 mmp-3="" pre="" recognition="" ribozyme="" site<="" ·=""></dd3></pre>	
	<400 + 4459	
ggcat	dagti destetatgg a	
	CL10 + 4460	
	0210 - 4400 0211 - 21	
	CIII > DNA	
	C113 - Homo sapien	
	The say I am a say I a	
	CML3 · MMP-3 ribozyme recognition site	
	400 · 4460	
cggaa	otgt coctocagaa c	2.
	<210 - 4461	
	<210 · 4461	
	K212 - DNA	
	:213 · Home sapien	
	4220 ·	
	<223 MMP-3 ribozyme recognition site	
	0400 + 4461	45.5
cctgc	ttgt citttgatge t	***
	F210 + 4462	
	221.	
	CALAN INA	
	0.113 · Home sapien	
	(220)	
	02230 MMP-3 ribozyme recognition site	
	(400): 4463	
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ccyac		4
	00100-4463	
	00100-4463 00110-71	
	00100 PNA	
	00130 Homo sapien	
	20 MA	
	<pre>COMMO: COMMP-9 ribozyme recognition site</pre>	
	or man=9 ribozyme recognition site	
	14001-14463	
agece	tagt cotigatgete e	21
	12101-14464	
	III. DE DIA	
	(213> Homo sapien	

	:220 > :223 > MMP-9 ribozyme recognition site	
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	H210 + 4465 H211 > 21 H212 > DNA H213 > Homo sapien	
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•	<210> 4466 <211> 21 <212> DNA <213> Homo sapien	
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	<220 > <223> MMP-9 ribozyme recognition site	
	८४००५ ४४६७ व्यक्ष occagacotg g	21
· · · · · · · · · · · · · · · · · · ·	<pre>&lt;3108 4468 &lt;1118 01 &lt;1118 DNA &lt;1118 Homo sapien</pre>	
	02200 02230 MMP-9 ribozyme recognition site	
	04005 4468 togt catocagttt g	21
	1210: 4469 1211: 21 1217: LMA 1213: Homo sapien	
•	(220)	

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-0023 · MMP-9 ribozyme	e recognition	site	
ા400 - 4469 agtttgqtqt દયદદુયagcac g			21
<pre>+0010 + 4470 +0011 + 01 +0012 + 5NA +0013 + Home sapien</pre>			
+220+ +223× MMP-9 ribozyme	e recognition	site	
-400× 4470 gagttgrygt bestaggeaa g			21
+210 × 4471 +211 + 21 +212 + DNA +213 × Homo sapien			
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-00200 -00230 MMP-9 ribozyme	e recognition	site	
k400% 447% caccgacgut cqctccgacg g			21
<pre>SC100 4473 +C110 C1 +C110 DNA +C130 Home sapien</pre>			
k.∟wk klij3k MMP-9 ribozyme	e recognition	site	
04000-4475 caeggaeggt egeteegaeg g			21
<pre>+0010: 4474 +0011: 01 +0210: DNA +0213: Homo sapien</pre>			
<2210 4223> MMP-9 ribozyme	e recognition	site	

ार्थणाः 4474 agetgtgogt etteseette a	.21
<pre>#1210 + 4475 #1211 + 21 #1212 + DNA #1213 + Homo sapien</pre>	
<pre>+MAG+ +MAP+9 ribozyme recognition site</pre>	
$\approx 4400 + 4475$ obtotatiggt bottogoodig a	21
<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	
<pre>&lt;0.00 + &lt;0.03 MMP-9 ribozyme recognition site</pre>	
04009-4476 dedegaaggt stigdeedadd g	21
<pre>%210 * 4477 %211 * 21 %212 * DNA %213 * Homo sapien</pre>	
<pre>&lt;220 + &lt;223 * MMP-9 ribozyme recognition site</pre>	
-400 + 4477 cocceatigt coaccectea g	21
<pre>&lt;:210&gt; 4478 &lt;:211&gt; 21 &lt;:112</pre>	
<pre><dre><dre><dre><dre><dre><dre><dre><d< td=""><td></td></d<></dre></dre></dre></dre></dre></dre></dre></pre>	
$\pm (400) \pm 4478$ gcotttqaqt coqgtggacg a	21
+1210+ 4479 +1211+ 21 +1213+ DMA +1213+ Homo sapien	
<pre>-(2000)(2007)- MMP-9 ribozyme recognition site</pre>	
:400: 4479 tggactoggt otttgaggag o	21

<pre>%010 + 4480 %011 + 21 %212 + DNA %213 + Homo sapien</pre>	
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<pre>-(210)+ (213)+ FDGF A ribozyme recognition site</pre>	
(400) 448) deggeoggit egeteetgaa g	21
-00100-4484 -0010-01 -00100-EMA -00130-Homo sapier	
-CARO	
H400: 4484 coccttong: coccaccoc a	21
.:21m. 4485	

.:211 •	21	
-1212 -		
	Homo sapien	
	nemo bapion	
+:22p+		
	PDGF A ribozyme recognition site	
-1400-	4485	
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cygycy, .	3.430ag000 0	• •
-1210 -	4486	
	ONA	
	Homo sapien	
-:220 -		
	FOGF A ribozyme recognition site	
-1400 ×	4486	
ggcccgcagt c		2:
<210 >	4487	
·311>	21	
<2212 €		
-:213 →	Homo sapien	
<12.20 ×		
+12.2.3 %	PDGF A ribozyme recognition site	
+(4 () () ×		
ctcacgaggi c	ccatgocact a	
-:210 ·		
-1211 -		
-:212	EMA	
KI213.5	Homo sapien	
W. V. V.		
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1 4 4 5	PDGF A ribozyme recognition site	
1.0.0	11115	
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	Homo sapien	
ئى 1 يەر <sup>ئ</sup>	TRANS SAPIET	
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	FDGF A ribozyme recognition site	
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1813 - Homo sapie	[1]		
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+12.20+ +12.23+ POOF A rib	ozyme recognition si	ite	
-400 - 4491 tootogdagt caddtogado	c		21
<pre>&lt;3210 / 4492 &lt;211 / 21 &lt;212 / DNA &lt;213 / Homo sapie:</pre>	n		
<pre></pre>	ozyme recognition si	ite	
k400 % 4492 ogagoagtgt baagtgobag (	c		21
<2105 4493 <211> 21 <2125 ENA <2135 Homo sapie:	n		
+2205 <223> FDGF A rib	ozyme recognition si	ite	
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<pre>&lt;0.10&gt; 44.34 &lt;0.11&gt; 0.1 &lt;0.10&gt; DNA &lt;0.15&gt; Homo sapier</pre>	n		
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+0.100+4495 +0.110+0.1 +0.110+DMA +0.2130+Hcmo-sapier	٦		

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+:220 ÷	GF B rikozyme	recognition	site	
-(400 - 44) aggttggagt cocc				21
-02100- 450 -02110- 21 -02120- DMA -02130- Hem	ı.			
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-:400+ 4500 ggcccgmagt cagpatgaat c	.?1
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<pre>&lt;120.* &lt;123. PDGP B ribozyme recognition site</pre>	
+4400 + 4501 ctottootgt ctottootgt c	.21
<pre>&lt;210 + 4502 &lt;211 + 21 &lt;212 + DNA &lt;213 &gt; Home sapien</pre>	
<220> <223> PDGF B ribozyme recognition site	
04008-4502 ctacctgpqt ctggtcagcg c	21
<210 > 4503 <211 > 21 <212 · DNA <213 > Homo sapien <220 ·	
-:223 - FTGF B ribozyme recognition site	
+400 + 4505 tgagtatqqt caqaqaqag g	::1
<pre>&lt;210&gt; 4504 <c11> D1 <c10> TMA <c13> Homo sapien</c13></c10></c11></pre>	
-:220s -:223- PDGF B ribozyme recognition site	
+400+4504 tgcgacciqi ccaqqtgaga a	21
+0.100+4505 +0.110+21 +0.100+DNA +0.130+Homo sapien	
<pre>(0.7()) &lt;2.33  PDGF B ribozyme recognition site</pre>	
€400 € 4505	

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<pre>0010 + 4506 0011 + 01 0011 + DNA 0013 + Homo sapien</pre>	
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<pre>&lt;:210 &gt; 4508 ::211 &gt; 11 ::212 &gt; DNA ::213 + Homo sapien</pre>	
<pre>&lt;220. &lt;223. PDGF B riboxyme recognition site</pre>	
(400% 450% gtoogtotgt otngatgoot g	21
0210 + 4509 0311 + 21 0310 + DNA 0313 > Hemo sapien	
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<pre>HDD08 HDGF B ribozyme recognition site</pre>	
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(213) VEGF hammerhead ribozyme recognition site	
1406> 4521 agaaaaaaaa toagttogag g	
1210> 4522 1211> 19 1212> DNA 1213> Homo sapien	21
<pre>/320. /323/ VEGF hammerhead ribozyme recognition site</pre>	
1400% 4522 gcaagaaato poggtataa	
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	19